FEB , 8 7005 E

SEQUENCE LISTING

```
<110> Bell, John C.
      Sonenberg, Nahum
      Stojdl, David F.
      Brown, Earl G.
      Atkins, Harold L.
      Marius, Ricardo M.
      Lichty, Brian D.
      Knowles, Shane B.
<120> ONCOLYTIC VIRUS
<130> 18003
<140> US 09/664,444
<141> 2000-09-18
<150> 60/287,590
<151> 1999-09-17
<160> 52
<170> PatentIn Ver. 2.1
<210> 1
<211> 11
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      peptide
Cys Thr Thr His Arg His His Thr Ser Asn Cys
                  5
<210> 2
<211> 12
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      peptide
<400> 2
Cys Leu Asn Ala His Arg Thr Thr His His Cys
                  5
<210> 3
<211> 11
<212> PRT
<213> Artificial Sequence
```

```
<220>
<223> Description of Artificial Sequence: Synthetic
     peptide
<400> 3
Cys His Gly Leu His Ser Asn Met Arg His Cys
         5
<210> 4
<211> 9
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     peptide
<400> 4
Cys His His His Arg Leu Asn Cys
                 5
<210> 5
<211> 9
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     peptide
<400> 5
Cys His Ser His His His Arg Gly Cys
 1
       5
<210> 6
<211> 9
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     peptide
<400> 6
Cys Trp Asp His His Asn His His Cys
 1
                 5
<210> 7
<211> 9
<212> PRT
<213> Artificial Sequence
```

```
<220>
<223> Description of Artificial Sequence: Synthetic
      peptide
<400> 7
Cys Asp Asn Asn His His His Cys
<210> 8
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     peptide
<400> 8
Cys His His Arg Ile Ser Ser His Cys
                  5
<210> 9
<211> 1269
<212> DNA
<213> Vesicular stomatitis virus
<400> 9
atgtctgtta cagtcaagag aatcattgac aacacagtca tagttccaaa acttcctgca 60
aatgaggatc cagtggaata cccggcagat tacttcagaa aatcaaagga gattcctctt 120
tacatcaata ctacaaaaag tttgtcagat ctaagaggat atgtctacca aggcctcaaa 180
tccggaaatg tatcaatcat acatgtcaac agctacttgt atggagcatt aaaggacatc 240
cggggtaagt tggataaaga ttggtcaagt ttcggaataa acatcgggaa agcaggggat 300
acaatcggaa tatttgacct tgtatccttg aaagccctgg acggcgtact tccagatgga 360
gtatcggatg cttccagaac cagcgcagat gacaaatggt tgcctttgta tctacttggc 420
ttatacagag tgggcagaac acaaatgcct gaatacagaa aaaagctcat ggatgggctg 480
acaaatcaat gcaaaatgat caatgaacag tttgaacctc ttgtgccaga aggtcgtgac 540
atttttgatg tgtggggaaa tgacagtaat tacacaaaaa ttgtcgctgc agtggacatg 600
ttcttccaca tgttcaaaaa acatgaatgt gcctcgttca gatacggaac tattgtttcc 660
agattcaaag attgtgctgc attggcaaca tttggacacc tctgcaaaat aaccggaatg 720
tctacagaag atgtaacgac ctggatcttg aaccgagaag ttgcagatga aatggtccaa 780
atgatgcttc caggccaaga aattgacaag gccgattcat acatgcctta tttgatcgac 840
tttggattgt cttctaagtc tccatattct tccgtcaaaa accctgcctt ccacttctgg 900
gggcaattga cagctcttct gctcagatcc accagagcaa ggaatgcccg acagcctgat 960
gacattgagt atacatctct tactacagca ggtttgttgt acgcttatgc agtaggatcc 1020
tctgccgact tggcacaaca gttttgtgtt ggagataaca aatacactcc agatgatagt 1080
accggaggat tgacgactaa tgcaccgcca caaggcagag atgtggtcga atggctcgga 1140
tggtttgaag atcaaaacag aaaaccgact cctgatatga tgcagtatgc gaaaagagca 1200
gtcatgtcac tgcaaggcct aagagagaag acaattggca agtatgctaa gtcagaattt 1260
                                                                  1269
gacaaatga
<210> 10
<211> 1269
<212> DNA
<213> Vesicular stomatitis virus
```

<400> 10 atgtctgtta cagtcaagag aatcattgcc aacacagtca tagttccaaa acttcctgca 60 aatgaggatc cagtggaata cccggcagat tacttcagaa aatcaaagga gattcctctt 120 tacatcaata ctacaaaaag tttgtcagat ctaagaggat atgtctacca aggcctcaaa 180 tccggaaatg tatcaatcat acatgtcaac agctacttgt atggagcatt gaaggacatc 240 cggggtaagt tggataaaga ttggtcaagt ttcggaataa acatcgggaa ggcaggggat 300 acaatcggaa tatttgacct tgtatccttg aaagccctgg acggtgtact tccagatgga 360 gtatcggatg cttccagaac cagcgcagat gacaaatggt tgcctttgta tctacttggc 420 ttatacagag tgggcagaac acaaatgcct gaatacagaa aaaggctcat ggatgggctg 480 acaaatcaat gcaaaatgat caatgaacag tttgaacctc ttgtgccaga aggtcgtgac 540 atttttgatg tgtggggaaa tgacagtaat tacacaaaaa ttgtcgctgc agtggacatg 600 ttcttccaca tgttcaaaaa acatgaatgt gcctcgttca gatacggaac tattgtttcc 660 agattcaaag attgtgctgc attggcaaca tttggacacc tctgcaaaat aaccggaatg 720 tctacagaag atgtaacgac ctggatcttg aaccgagaag ttgcagatga gatggtccaa 780 atgatgette caggecaaga aattgacaag geegatteat acatgeetta tttgategae 840 tttggattgt cttctaagtc tccatattct tccgtcaaaa accctgcctt ccacttctgg 900 gggcaattga cagctcttct gctcagatcc accagagcaa ggaatgcccg acagcctgat 960 gacattgagt atacatctct tactacagca ggtttgttgt acgcttatgc agtaggatcc 1020 tctgctgact tggcacaaca gttttgtgtt ggagatagca aatacactcc agatgatagt 1080 accggaggat tgacgactaa tgcaccgcca caaggcagag atgtggtcga atggctcgga 1140 tggtttgaag atcaaaacag aaaaccgact cctgatatga tgcagtatgc gaaacgagca 1200 gtcatgtcac tgcaaggcct aagagagaag acaattggca agtatgctaa gtcagaattt 1260 1269 gacaaatga <210> 11 <211> 1269 <212> DNA <213> Vesicular stomatitis virus <400> 11 atgtctgtta cagtcaagag aatcattgac aacacagtca tagttccaaa acttcctgca 60 aatgaggatc cagtggaata cccggcagat tacttcagaa aatcaaagga gattcctctt 120 tacatcaata ctacaaaaag tttgtcagat ctaagaggat atgtctacca aggcctcaaa 180 tccggaaatg tatcaatcat acatgtcaac agctacttgt atggagcatt gaaggacatc 240 cggggtaagt tggataaaga ttggtcaagt ttcggaataa acatcgggaa ggcaggggat 300 acaatcggaa tatttgacct tgtatccttg aaagccctgg acggtgtact tccagatgga 360 gtatcggatg cttccagaac cagcgcagat gacaaatggt tgcctttgta tctacttggc 420 ttatacagag tgggcagaac acaaatgcct gaatacagaa aaaggctcat ggatgggctg 480 acaaatcaat gcaaaatgat caatgaacag tttgaacctc ttgtgccaga aggtcgtgac 540 atttttgatg tgtggggaaa tgacagtaat tacacaaaaa ttgtcgctgc agtggacatg 600 ttcttccaca tgttcaaaaa acatgaatgt gcctcgttca gatacggaac tattgtttcc 660 agattcaaag attgtgctgc attggcaaca tttggacacc tctgcaaaat aaccggaatg 720 tctacagaag atgtaacgac ctggatcttg aaccgagaag ttgcagatga gatggtccaa 780 atgatgette caggecaaga aattgacaag geegatteat acatgeetta tttgategae 840 tttggattgt cttctaagtc tccatattct tccgtcaaaa accctgcctt ccacttctgg 900 gggcaattga cagctcttct gctcagatct accagagcaa ggaatgcccg acagcctgat 960 gacattgagt atacatetet tactacagea ggtttgttgt acgettatge agtaggatee 1020 tctgctgact tggcacaaca gttttgtgtt ggagatagca aatacactcc agatgatagt 1080 accggaggat tgacgactaa tgcaccgcca caaggcagag atgtggtcga atggctcgga 1140 tggtttgaag atcaaaacag aaaaccgact cctgatatga tgcagtatgc gaaacgagca 1200 gtcatgtcac tgcaaggcct aagagagaag acaattggca agtatgctaa gtcagaattt 1260

1269

gacaaatga

```
<211> 1269
<212> DNA
<213> Vesicular stomatitis virus
<400> 12
atgtctgtta cagtcaagag aatcattgac aacacagtca tagttccaaa acttcctgca 60
aatgaggatc cagtggaata cccggcagat tacttcagaa aatcaaagga gattcctctt 120
tacatcaata ctacaaaaag tttgtcagat ctaagaggat atgtctacca aggcctcaaa 180
tccggaaatg tatcaatcat acatgtcaac agctacttgt atggagcatt gaaggacatc 240
cggggtaagt tggataaaga ttggtcaagt ttcggaataa acatcgggaa ggcaggggat 300
acaateggaa tatttgacet tgtateettg aaageeetgg aeggtgtaet teeagatgga 360
gtatcggatg cttccagaac cagcgcagat gacaaatggt tgcctttgta tctacttggc 420
ttatacagag tgggcagaac acaaatgcct gaatacagaa aaaggctcat ggatgggctg 480
acaaatcaat gcaaaatgat caatgaacag tttgaacctc ttgtgccaga aggtcgtgac 540
atttttgatg tgtggggaaa tgacagtaat tacacaaaaa ttgtcgctgc agtggacatg 600
ttcttccaca tgttcaaaaa acatgaatgt gcctcgttca gatacggaac tattgtttcc 660
agattcaaag attgtgctgc attggcaaca tttggacacc tctgcaaaat aaccggaatg 720
tctacagaag atgtaacgac ctggatcttg aaccgagaag ttgcagatga gatggtccaa 780
atgatgcttc caggccaaga aattgacaag gccgattcat acatgcctta tttgatcgac 840
tttggattgt cttctaagtc tccatattct tccgtcaaaa accctgcctt ccacttctgg 900
gggcaattga cagetettet geteagatee accagageaa ggaatgeeeg acageetgat 960
gacattgagt atacatctct tactacagca ggtttgttgt acgcttatgc agtaggatcc 1020
tctgctgact tggcacaaca gttttgtgtt ggagatagca aatacactcc agatgatagt 1080
accggaggat tgacgactaa tgcaccgcca caaggcagag atgtggtcga atggctcgga 1140
tggtttgaag atcaaaacag aaaaccgact cctgatatga tgcagtatgc gaaacgagca 1200
gtcatgtcac tgcaaggcct aagagagaag acaattggca agtatgctaa gtcagaattt 1260
gacaaatga
<210> 13
<211> 1029
<212> DNA
<213> Vesicular stomatitis virus
<220>
<221> modified_base
<222> (1)..(1029)
<223> "n" represents a, t, c, g, other or unknown
<400> 13
tcaatcatac atgtcaacag ctacttgtat ggagcattga aggacatccg gggtaagttg 60
gataaagatt ggtcaagttt cggaataaac atcgggaagg caggggatac aatcggaata 120
tttgaccttg tatccttgaa agccctggac ggtgtacttc cagatggagt atcggatgct 180
tccagaacca gcgcagatga caaatggttg cctttgtatc tacttggctt atacagagtg 240
ggcagaacac aaatgcctga atacagaaaa aggctcatgg atgggctgac aaatcaatgc 300
aaaatgatca atgaacagtt tgaacctctt gtgccagaag gtcgtgacat ttttgatgtg 360
tggggaaatg acagtaatta cacaaaaatt gtcgctgcag tggacatgtt cttccacatg 420
ttcaaaaaac atgaatgtgc ctcgttcaga tacggaacta ttgtttccag attcaaagat 480
tgtgctgcat tggcaacatt tggacacctc tgcaaaataa ccggaatgtc tacagaagat 540
gtaacgacct ggatcttgaa ccgagaagtt gcagatgaga tggtccaaat gatgcttcca 600
ggccaagaaa ttgacaaggc cgattcatac atgccttatt tgatcgactt tggattgtct 660
tctaagtctc catattcttc cgtcaaaaac cctgccttcc acttctgggg gcaattgact 720
gacattgagt atacateten tactacagea ggtttgttgt acgettatge agtaggatee 780
tetgetgact tggcacanca gttttgtgtt ggagatagca aatacactcc agatgatagt 840
accggaggat tgacgactaa tgcaccgcca caaggcagag atgtggtcga atggctcgga 900
tggtttgaag atcaaaacag aaaaccgact cctgatatga tgcagtatgc gaaacgagca 960
gtcatgtcac tgcaaggcct aagagagaag acaattggca agtatgctaa gtcagaattt 1020
```

gacaaatga 1029

<213 <213)> 14 L> 42 2> PI B> Ve	22 RT	ular	stor	mati	tis v	virus	5							
)> 14 Ser		Thr	Val 5	Lys	Arg	Ile	Ile	Asp 10	Asn	Thr	Val	Ile	Val 15	Pro
Lys	Leu	Pro	Ala 20	Asn	Glu	Asp	Pro	Val 25	Glu	Tyr	Pro	Ala	Asp 30	Tyr	Phe
Arg	Lys	Ser 35	Lys	Glu	Ile	Pro	Leu 40	Tyr	Ile	Asn	Thr	Thr 45	Lys	Ser	Leu
Ser	Asp 50	Leu	Arg	Gly	Tyr	Val 55	Tyr	Gln	Gly	Leu	Lys 60	Ser	Gly	Asn	Val
Ser 65	Ile	Ile	His	Val	Asn 70	Ser	Tyr	Leu	Tyr	Gly 75	Ala	Leu	Lys	Asp	Ile 80
Arg	Gly	Lys	Leu	Asp 85	Lys	Asp	Trp	Ser	Ser 90	Phe	Gly	Ile	Asn	Ile 95	Gly
Lys	Ala	Gly	Asp 100	Thr	Ile	Gly	Ile	Phe 105	Asp	Leu	Val	Ser	Leu 110	Lys	Ala
Leu	Asp	Gly 115	Val	Leu	Pro	Asp	Gly 120	Val	Ser	Asp	Ala	Ser 125	Arg	Thr	Ser
Ala	Asp 130	Asp	Lys	Trp	Leu	Pro 135	Leu	Tyr	Leu	Leu	Gly 140	Leu	Tyr	Arg	Val
Gly 145	Arg	Thr	Gln	Met	Pro 150	Glu	Tyr	Arg	Lys	Lys 155	Leu	Męt	Asp	Gly	Leu 160
Thr	Asn	Gln	Cys	Lys 165	Met	Ile	Asn	Glu	Gln 170	Phe	Glu	Pro	Leu	Val 175	Pro
Glu	Gly	Arg	Asp 180	Ile	Phe	Asp	Val	Trp 185	Gly	Asn	Asp	Ser	Asn 190	Tyr	Thr
Lys	Ile	Val 195	Ala	Ala	Val	Asp	Met 200	Phe	Phe	His	Met	Phe 205	Lys	Lys	His
Glu	Cys 210	Ala	Ser	Phe	Arg	Tyr 215	Gly	Thr	Ile	Val	Ser 220	Arg	Phe	Lys	Asp

Cys Ala Ala Leu Ala Thr Phe Gly His Leu Cys Lys Ile Thr Gly Met

230 . 235

Ser Thr Glu Asp Val Thr Thr Trp Ile Leu Asn Arg Glu Val Ala Asp

250

255

245

225

Glu Met Val Gln Met Met Leu Pro Gly Gln Glu Ile Asp Lys Ala Asp 265 260 Ser Tyr Met Pro Tyr Leu Ile Asp Phe Gly Leu Ser Ser Lys Ser Pro 280 Tyr Ser Ser Val Lys Asn Pro Ala Phe His Phe Trp Gly Gln Leu Thr 295 Ala Leu Leu Leu Arg Ser Thr Arg Ala Arg Asn Ala Arg Gln Pro Asp 310 Asp Ile Glu Tyr Thr Ser Leu Thr Thr Ala Gly Leu Leu Tyr Ala Tyr 330 Ala Val Gly Ser Ser Ala Asp Leu Ala Gln Gln Phe Cys Val Gly Asp 340 345 Asn Lys Tyr Thr Pro Asp Asp Ser Thr Gly Gly Leu Thr Thr Asn Ala 360 Pro Pro Gln Gly Arg Asp Val Val Glu Trp Leu Gly Trp Phe Glu Asp 375 Gln Asn Arg Lys Pro Thr Pro Asp Met Met Gln Tyr Ala Lys Arg Ala 390 385 Val Met Ser Leu Gln Gly Leu Arg Glu Lys Thr Ile Gly Lys Tyr Ala 410 Lys Ser Glu Phe Asp Lys 420 <210> 15 <211> 422 <212> PRT <213> Vesicular stomatitis virus <400> 15 Met Ser Val Thr Val Lys Arg Ile Ile Ala Asn Thr Val Ile Val Pro Lys Leu Pro Ala Asn Glu Asp Pro Val Glu Tyr Pro Ala Asp Tyr Phe 30 20 Arg Lys Ser Lys Glu Ile Pro Leu Tyr Ile Asn Thr Thr Lys Ser Leu Ser Asp Leu Arg Gly Tyr Val Tyr Gln Gly Leu Lys Ser Gly Asn Val 55 Ser Ile Ile His Val Asn Ser Tyr Leu Tyr Gly Ala Leu Lys Asp Ile 70 65

Arg Gly Lys Leu Asp Lys Asp Trp Ser Ser Phe Gly Ile Asn Ile Gly

90

85

Lys Ala Gly	Asp Thr I	e Gly Ile	Phe Asp	Leu Val	Ser Leu 110	Lys Ala
Leu Asp Gly 115	Val Leu P	o Asp Gly 120		Asp Ala	Ser Arg 125	Thr Ser
Ala Asp Asp 130	Lys Trp L	eu Pro Leu 135	Tyr Leu	Leu Gly 140	Leu Tyr	Arg Val
Gly Arg Thr 145		ro Glu Tyr 50	Arg Lys	Arg Leu 155	Met Asp	Gly Leu 160
Thr Asn Gln	Cys Lys Mo 165	et Ile Asn	Glu Gln 170	Phe Glu	Pro Leu	Val Pro 175
Glu Gly Arg	Asp Ile Pl 180	ne Asp Val	Trp Gly 185	Asn Asp	Ser Asn 190	Tyr Thr
Lys Ile Val 195	Ala Ala V	al Asp Met 200		His Met	Phe Lys 205	Lys His
Glu Cys Ala 210	Ser Phe A	g Tyr Gly 215	Thr Ile	Val Ser 220	Arg Phe	Lys Asp
Cys Ala Ala 225		nr Phe Gly 30	His Leu	Cys Lys 235	Ile Thr	Gly Met 240
Ser Thr Glu	Asp Val Ti 245	nr Thr Trp	Ile Leu 250	Asn Arg	Glu Val	Ala Asp 255
Glu Met Val	Gln Met Me 260	et Leu Pro	Gly Gln 265	Glu Ile	Asp Lys 270	Ala Asp
Ser Tyr Met 275	Pro Tyr L	eu Ile Asp 280	_	Leu Ser	Ser Lys 285	Ser Pro
Tyr Ser Ser 290	Val Lys A	sn Pro Ala 295	Phe His	Phe Trp 300	Gly Gln	Leu Thr
Ala Leu Leu 305	_	er Thr Arg 10	Ala Arg	Asn Ala 315	Arg Gln	Pro Asp 320
Asp Ile Glu	Tyr Thr So	er Leu Thr	Thr Ala	Gly Leu	Leu Tyr	Ala Tyr 335
Ala Val Gly	Ser Ser A	la Asp Lev	Ala Gln 345	Gln Phe	Cys Val 350	Gly Asp
Ser Lys Tyr 355	Thr Pro A	sp Asp Ser 360	_	Gly Leu	Thr Thr	Asn Ala
Pro Pro Gln 370	Gly Arg A	sp Val Val 375	Glu Trp	Leu Gly 380	Trp Phe	Glu Asp
Gln Asn Arg 385		nr Pro Asp 90	Met Met	Gln Tyr 395	Ala Lys	Arg Ala 400

Val Met Ser Leu Gln Gly Leu Arg Glu Lys Thr Ile Gly Lys Tyr Ala 405 410 415

Lys Ser Glu Phe Asp Lys 420

<210> 16

<211> 422

<212> PRT

<213> Vesicular stomatitis virus

<400> 16

Met Ser Val Thr Val Lys Arg Ile Ile Asp Asn Thr Val Ile Val Pro 1 5 10 15

Lys Leu Pro Ala Asn Glu Asp Pro Val Glu Tyr Pro Ala Asp Tyr Phe 20 25 30

Arg Lys Ser Lys Glu Ile Pro Leu Tyr Ile Asn Thr Thr Lys Ser Leu 35 40 45

Ser Asp Leu Arg Gly Tyr Val Tyr Gln Gly Leu Lys Ser Gly Asn Val 50 55 60

Ser Ile Ile His Val Asn Ser Tyr Leu Tyr Gly Ala Leu Lys Asp Ile 65 70 75 80

Arg Gly Lys Leu Asp Lys Asp Trp Ser Ser Phe Gly Ile Asn Ile Gly 85 90 95

Lys Ala Gly Asp Thr Ile Gly Ile Phe Asp Leu Val Ser Leu Lys Ala 100 105 110

Leu Asp Gly Val Leu Pro Asp Gly Val Ser Asp Ala Ser Arg Thr Ser 115 120 125

Ala Asp Asp Lys Trp Leu Pro Leu Tyr Leu Leu Gly Leu Tyr Arg Val 130 135 140

Gly Arg Thr Gln Met Pro Glu Tyr Arg Lys Arg Leu Met Asp Gly Leu 145 150 155 160

Thr Asn Gln Cys Lys Met Ile Asn Glu Gln Phe Glu Pro Leu Val Pro 165 170 175

Glu Gly Arg Asp Ile Phe Asp Val Trp Gly Asn Asp Ser Asn Tyr Thr 180 185 190

Lys Ile Val Ala Ala Val Asp Met Phe Phe His Met Phe Lys Lys His
195 200 205

Glu Cys Ala Ser Phe Arg Tyr Gly Thr Ile Val Ser Arg Phe Lys Asp 210 215 220

Cys Ala Ala Leu Ala Thr Phe Gly His Leu Cys Lys Ile Thr Gly Met

Ser Thr Glu Asp Val Thr Trp Ile Leu Asn Arg Glu Val Ala Asp 245 250 255

Glu Met Val Gln Met Met Leu Pro Gly Gln Glu Ile Asp Lys Ala Asp 260 265 270

Ser Tyr Met Pro Tyr Leu Ile Asp Phe Gly Leu Ser Ser Lys Ser Pro 275 280 285

Tyr Ser Ser Val Lys Asn Pro Ala Phe His Phe Trp Gly Gln Leu Thr 290 295 300

Ala Leu Leu Leu Arg Ser Thr Arg Ala Arg Asn Ala Arg Gln Pro Asp 305 310 315 320

Asp Ile Glu Tyr Thr Ser Leu Thr Thr Ala Gly Leu Leu Tyr Ala Tyr 325 330 335

Ala Val Gly Ser Ser Ala Asp Leu Ala Gln Gln Phe Cys Val Gly Asp 340 345 350

Ser Lys Tyr Thr Pro Asp Asp Ser Thr Gly Gly Leu Thr Thr Asn Ala 355 360 365

Pro Pro Gln Gly Arg Asp Val Val Glu Trp Leu Gly Trp Phe Glu Asp 370 375 380

Gln Asn Arg Lys Pro Thr Pro Asp Met Met Gln Tyr Ala Lys Arg Ala 385 390 395 400

Val Met Ser Leu Gln Gly Leu Arg Glu Lys Thr Ile Gly Lys Tyr Ala 405 410 415

Lys Ser Glu Phe Asp Lys 420

<210> 17

<211> 342

<212> PRT

<213> Vesicular stomatitis virus

<220>

<221> MOD_RES

<222> (1)..(342)

<223> "Xaa" represents any, other or unknown amino acid

<400> 17

Ser Ile Ile His Val Asn Ser Tyr Leu Tyr Gly Ala Leu Lys Asp Ile 1 5 10 15

Arg Gly Lys Leu Asp Lys Asp Trp Ser Ser Phe Gly Ile Asn Ile Gly 20 25 30

Lys Ala Gly Asp Thr Ile Gly Ile Phe Asp Leu Val Ser Leu Lys Ala

Lys Ser Glu Phe Asp Lys

Leu	Asp 50	Gly	Val	Leu	Pro	Asp 55	Gly	Val	Ser	Asp	Ala 60	Ser	Arg	Thr	Ser
Ala 65	Asp	Asp	Lys	Trp	Leu 70	Pro	Leu	Tyr	Leu	Leu 75	Gly	Leu	Tyr	Arg	Va] 80
Gly	Arg	Thr	Gln	Met 85	Pro	Glu	Tyr	Arg	Lys 90	Arg	Leu	Met	Asp	Gly 95	Leu
Thr	Asn	Gln	Cys 100	Lys	Met	Ile	Asn	Glu 105	Gln	Phe	Glu	Pro	Leu 110	Val	Pro
Glu	Gly	Arg 115	Asp	Ile	Phe	Asp	Val 120	Trp	Gly	Asn	Asp	Ser 125	Asn	Tyr	Thr
Lys	Ile 130	Val	Ala	Ala	Val	Asp 135	Met	Phe	Phe	His	Met 140	Phe	Lys	Lys	His
Glu 145	Cys	Ala	Ser	Phe	Arg 150	Tyr	Gly	Thr	Ile	Val 155	Ser	Arg	Phe	Lys	Asp 160
Cys	Ala	Ala	Leu	Ala 165	Thr	Phe	Gly	His	Leu 170	Cys	Lys	Ile	Thr	Gly 175	Met
Ser	Thr	Glu	Asp 180	Val	Thr	Thr	Trp	Ile 185	Leu	Asn	Arg	Glu	Val 190	Ala	Asp
Glu	Met	Val 195	Gln	Met	Met	Leu	Pro 200	Gly	Gln	Glu	Ile	Asp 205	Lys	Ala	Asp
Ser	Tyr 210	Met	Pro	Tyr	Leu	Ile 215	Asp	Phe	Gly	Leu	Ser 220	Ser	Lys	Ser	Pro
Tyr 225	Ser	Ser	Val	Lys	Asn 230	Pro	Ala	Phe	His	Phe 235	Trp	Gly	Gln	Leu	Thr 240
Asp	Ile	Glu	Tyr	Thr 245	Ser	Xaa	Thr	Thr	Ala 250	Gly	Leu	Leu	Tyr	Ala 255	Tyr
Ala	Val	Gly	Ser 260	Ser	Ala	Asp	Leu	Ala 265	Xaa	Gln	Phe	Cys	Val 270	Gly	Asp
Ser	Lys	Tyr 275	Thr	Pro	Asp	Asp	Ser 280	Thr	Gly	Gly	Leu	Thr 285	Thr	Asn	Ala
Pro	Pro 290	Gln	Gly	Arg	Asp	Val 295	Val	Glu	Trp	Leu	Gly 300	Trp	Phe	Glu	Asp
Gln 305	Asn	Arg	Lys	Pro	Thr 310	Pro	Asp	Met	Met	Gln 315	Tyr	Ala	Lys	Arg	Ala 320
Val	Met	Ser	Leu	Gln 325	Gly	Leu	Arg	Glu	Lys 330	Thr	Ile	Gly	Lys	Tyr 335	Ala

```
<210> 18
<211> 798
<212> DNA
<213> Vesicular stomatitis virus
<400> 18
atggataatc tcacaaaagt tcgtgagtat ctcaagtcct attctcgtct ggatcaggcg 60
gtaggagaga tagatgagat cgaagcacaa cgagctgaaa agtccaatta tgagttgttc 120
caagaggatg gagtggaaga gcatactaag ccctcttatt ttcaggcagc agatgattct 180
gacacagaat ctgaaccaga aattgaagac aatcaaggtt tgtatgcaca ggatccagaa 240
gctgagcaag ttgaaggctt tatacagggg cctttagatg actatgcaga tgaggaagtg 300
gatgttgtat ttacttcgga ctggaaacca cctgagcttg aatctgacga gcatggaaag 360
accttacggt tgacatcgcc agagggttta agtggagagc agaaatccca gtggctttcg 420
acgattaaag cagtcgtgca aagtgccaaa tactggaatc tggcagagtg cacatttgaa 480
gcatcgggag aaggggtcat tatgaaggag cgccagataa ctccggatgt atataaggtc 540
actocagtga tgaacacaca tccgtcccaa tcagaagcag tatcagatgt ttggtctctc 600
tcaaagacat ccatgacttt ccaacccaag aaagcaagtc ttcagcctct caccatatcc 660
ttggatgaat tgttctcatc tagaggagag ttcatctctg tcggaggtga cggacgaatg 720
tctcataaag aggccatcct gctcggcctg agatacaaaa agttgtacaa tcaggcgaga 780
                                                                  798
gtcaaatatt ctctgtag
<210> 19
<211> 798
<212> DNA
<213> Vesicular stomatitis virus
<400> 19
atggataatc tcacaaaagt tcgtgagtat ctcaagtcct attctcgtct agatcaggcg 60
gtaggagaga tagatgagat cgaagcacaa cgagctgaaa agtccaatta tgagttgttc 120
caagaggacg gagtggaaga gcatactagg ccctcttatt ttcaggcagc agatgattct 180
gacacagaat ctgaaccaga aattgaagac aatcaaggct tgtatgtacc agatccggaa 240
gctgagcaag ttgaaggctt tatacagggg cctttagatg actatgcgga tgaggacgtg 300
gatgttgtat tcacttcgga ctggaaacag cctgagcttg aatccgacga gcatggaaag 360
accttacggt tgacattgcc agagggttta agtggagagc agaaatccca gtggcttttg 420
acgattaaag cagtcgttca aagtgccaaa cactggaatc tggcagagtg cacatttgaa 480
gcatcgggag aaggggtcat cataaaaaag cgccagataa ctccggatgt atataaggtc 540
actccagtga tgaacacaca tccgtcccaa tcagaagccg tatcagatgt ttggtctctc 600
tcaaagacat ccatgacttt ccaacccaag aaagcaagtc ttcagcctct caccatatcc 660
ttggatgaat tgttctcatc tagaggagaa ttcatctctg tcggaggtaa cggacgaatg 720
tctcataaag aggccatcct gctcggtctg aggtacaaaa agttgtacaa tcaggcgaga 780
                                                                  798
gtcaaatatt ctctgtag
<210> 20
<211> 570
<212> DNA
<213> Vesicular stomatitis virus
<400> 20
atggataatc tcacaaaagt tcgtgagtat ctcaagtcct attctcgtct agatcaggcg 60
gtaggagaga tagatgagat cgaagcacaa cgagctgaaa agtccaatta tgagttgttc 120
caagaggacg gagtggaaga gcatactagg ccctcttatt ttcaggcagc agatgattct 180
gacacagaat ctgaaccaga aattgaagac aatcaaggct tgtatgtacc agatccggaa 240
```

```
gctgagcaag ttgaaggctt tatacagggg cctttagatg actatgcgga tgaggacgtg 300
gatgttgtat tcacttcgga ctggaaacag cctgagcttg aatccgacga gcatggaaag 360
accttacggt tgacattgcc agagggttta agtggagagc agaaatccca gtggcttttg 420
acgattaaag cagtcgttca aagtgccaaa cactggaatc tggcagagtg cacatttgaa 480
gcatcgggag aaggggtcat cataaaaaag cgccagataa ctccggatgt atataaggtc 540
                                                                  570
actccagtga tgaacacaca tccgtcccaa
<210> 21
<211> 798
<212> DNA
<213> Vesicular stomatitis virus
<400> 21
atggataatc tcacaaaagt tcgtgagtat ctcaagtcct attctcgtct agatcaggcg 60
gtaggagaga tagatgagat cgaagcacaa cgagctgaaa agtccaatta tgagttgttc 120
caagaggacg gagtggaaga gcatactagg ccctcttatt ttcaggcagc agatgattct 180
qacacagaat ctgaaccaga aattgaagac aatcaaggct tgtatgtacc agatccggaa 240
qctgagcaag ttgaaggctt tatacagggg cctttagatg actatgcgga tgaggacgtg 300
gatgttgtat tcacttcgga ctggaaacag cctgagcttg aatccgacga gcatggaaag 360
accttacggt tgacattgcc agagggttta agtggagagc agaaatccca gtggcttttg 420
acgattaaag cagtcgttca aagtgccaaa cactggaatc tggcagagtg cacatttgaa 480
gcatcgggag aaggggtcat cataaaaaag cgccagataa ctccggatgt atataaggtc 540
actocagtga tgaacacaca tocgtoccaa toggaagcog tatoagatgt ttggtototo 600
tcaaagacat ccatgacttt ccaacccaag aaagcaagtc ttcagcctct caccatatcc 660
ttggatgaat tgttctcatc tagaggagaa ttcatctctg tcggaggtaa cggacgaatg 720
tctcataaag aggccatcct gctcggtctg aggtacaaaa agttgtacaa tcaggcgaga 780
                                                                  798
gtcaaatatt ctctgtag
<210> 22
<211> 798
<212> DNA
<213> Vesicular stomatitis virus
<400> 22
atggataatc tcacaaaagt tcgtgagtat ctcaagtcct attctcgtct agatcaggcg 60
gtaggagaga tagatgagat cgaagcacaa cgagctgaaa agtccaatta tgagttgttc 120
caagaggacg gagtggaaga gcatactagg ccctcttatt ttcaggcagc agatgattct 180
gacacagaat ctgaaccaga aattgaagac aatcaaggct tgtatgtacc agatccggaa 240
gctgagcaag ttgaaggctt tatacagggg cctttagatg actatgcgga tgaggacgtg 300
gatgttgtat tcacttcgga ctggaaacag cctgagcttg aatccgacga gcatggaaag 360
accttacggt tgacattgcc agagggttta agtggagagc agaaatccca gtggcttttg 420
acqattaaaq caqtcqttca aaqtqccaaa cactgqaatc tggcagagtg cacatttgaa 480
gcatcgggag aaggggtcat cataaaaaag cgccagataa ctccggatgt atataaggtc 540
actocagtga tgaacacaca tccgtcccaa tcagaagccg tatcagatgt ttggtctctc 600
tcaaagacat ccatgacttt ccaacccaag aaagcaagtc ttcagcctct caccatatcc 660
ttggatgaat tgttctcatc tagaggagaa ttcatctctg tcggaggtaa cggacgaatg 720
tctcataaag aggccatcct gctcggtctg aggtacaaaa agttgtacaa tcaggcgaga 780
                                                                  798
gtcaaatatt ctctgtag
<210> 23
<211> 265
```

<212> PRT

<213> Vesicular stomatitis virus

<400> 23

Met Asp Asn Leu Thr Lys Val Arg Glu Tyr Leu Lys Ser Tyr Ser Arg

1 5 10 15

Leu Asp Gln Ala Val Gly Glu Ile Asp Glu Ile Glu Ala Gln Arg Ala 20 25 30

Glu Lys Ser Asn Tyr Glu Leu Phe Gln Glu Asp Gly Val Glu His
35 40 45

Thr Lys Pro Ser Tyr Phe Gln Ala Asp Asp Ser Asp Thr Glu Ser 50 55 60

Glu Pro Glu Ile Glu Asp Asn Gln Gly Leu Tyr Ala Gln Asp Pro Glu 65 70 75 80

Ala Glu Gln Val Glu Gly Phe Ile Gln Gly Pro Leu Asp Asp Tyr Ala 85 90 95

Asp Glu Glu Val Asp Val Val Phe Thr Ser Asp Trp Lys Pro Pro Glu
100 105 110

Leu Glu Ser Asp Glu His Gly Lys Thr Leu Arg Leu Thr Ser Pro Glu 115 120 125

Gly Leu Ser Gly Glu Gln Lys Ser Gln Trp Leu Ser Thr Ile Lys Ala 130 135 140

Val Val Gln Ser Ala Lys Tyr Trp Asn Leu Ala Glu Cys Thr Phe Glu 145 150 155 160

Ala Ser Gly Glu Gly Val Ile Met Lys Glu Arg Gln Ile Thr Pro Asp 165 170 175

Val Tyr Lys Val Thr Pro Val Met Asn Thr His Pro Ser Gln Ser Glu
180 185 190

Ala Val Ser Asp Val Trp Ser Leu Ser Lys Thr Ser Met Thr Phe Gln
195 200 205

Pro Lys Lys Ala Ser Leu Gln Pro Leu Thr Ile Ser Leu Asp Glu Leu 210 215 220

Phe Ser Ser Arg Gly Glu Phe Ile Ser Val Gly Gly Asp Gly Arg Met 225 230 235 240

Ser His Lys Glu Ala Ile Leu Leu Gly Leu Arg Tyr Lys Lys Leu Tyr 245 250 255

Asn Gln Ala Arg Val Lys Tyr Ser Leu 260 265

<210> 24

<211> 265

<212> PRT

<213> Vesicular stomatitis virus

<400> 24

Met Asp Asn Leu Thr Lys Val Arg Glu Tyr Leu Lys Ser Tyr Ser Arg

1 5 10 15

Leu Asp Gln Ala Val Gly Glu Ile Asp Glu Ile Glu Ala Gln Arg Ala 20 25 30

Glu Lys Ser Asn Tyr Glu Leu Phe Gln Glu Asp Gly Val Glu Glu His 35 40 45

Thr Arg Pro Ser Tyr Phe Gln Ala Ala Asp Asp Ser Asp Thr Glu Ser 50 55 60

Glu Pro Glu Ile Glu Asp Asn Gln Gly Leu Tyr Val Pro Asp Pro Glu 65 70 75 80

Ala Glu Gln Val Glu Gly Phe Ile Gln Gly Pro Leu Asp Asp Tyr Ala 85 90 95

Asp Glu Asp Val Asp Val Val Phe Thr Ser Asp Trp Lys Gln Pro Glu
100 105 110

Leu Glu Ser Asp Glu His Gly Lys Thr Leu Arg Leu Thr Leu Pro Glu 115 120 125

Gly Leu Ser Gly Glu Gln Lys Ser Gln Trp Leu Leu Thr Ile Lys Ala 130 135 140

Val Val Gln Ser Ala Lys His Trp Asn Leu Ala Glu Cys Thr Phe Glu 145 150 155 160

Ala Ser Gly Glu Gly Val Ile Ile Lys Lys Arg Gln Ile Thr Pro Asp 165 170 175

Val Tyr Lys Val Thr Pro Val Met Asn Thr His Pro Ser Gln Ser Glu 180 185 190

Ala Val Ser Asp Val Trp Ser Leu Ser Lys Thr Ser Met Thr Phe Gln
195 200 205

Pro Lys Lys Ala Ser Leu Gln Pro Leu Thr Ile Ser Leu Asp Glu Leu 210 215 220

Phe Ser Ser Arg Gly Glu Phe Ile Ser Val Gly Gly Asn Gly Arg Met 225 230 235 240

Ser His Lys Glu Ala Ile Leu Leu Gly Leu Arg Tyr Lys Lys Leu Tyr 245 250 255

Asn Gln Ala Arg Val Lys Tyr Ser Leu 260 265

<210> 25

<211> 190

<212> PRT

<213> Vesicular stomatitis virus

<400> 25

Met Asp Asn Leu Thr Lys Val Arg Glu Tyr Leu Lys Ser Tyr Ser Arg 1 5 10 15

Leu Asp Gln Ala Val Gly Glu Ile Asp Glu Ile Glu Ala Gln Arg Ala 20 25 30

Glu Lys Ser Asn Tyr Glu Leu Phe Gln Glu Asp Gly Val Glu Glu His
35 40 45

Thr Arg Pro Ser Tyr Phe Gln Ala Ala Asp Asp Ser Asp Thr Glu Ser 50 55 60

Glu Pro Glu Ile Glu Asp Asn Gln Gly Leu Tyr Val Pro Asp Pro Glu 65 70 75 80

Ala Glu Gln Val Glu Gly Phe Ile Gln Gly Pro Leu Asp Asp Tyr Ala 85 90 95

Asp Glu Asp Val Asp Val Val Phe Thr Ser Asp Trp Lys Gln Pro Glu
100 105 110

Leu Glu Ser Asp Glu His Gly Lys Thr Leu Arg Leu Thr Leu Pro Glu 115 120 125

Gly Leu Ser Gly Glu Gln Lys Ser Gln Trp Leu Leu Thr Ile Lys Ala 130 135 140

Val Val Gln Ser Ala Lys His Trp Asn Leu Ala Glu Cys Thr Phe Glu 145 150 155 160

Ala Ser Gly Glu Gly Val Ile Ile Lys Lys Arg Gln Ile Thr Pro Asp 165 170 175

Val Tyr Lys Val Thr Pro Val Met Asn Thr His Pro Ser Gln 180 185 190

<210> 26

<211> 265

<212> PRT

<213> Vesicular stomatitis virus

<400> 26

Met Asp Asn Leu Thr Lys Val Arg Glu Tyr Leu Lys Ser Tyr Ser Arg
1 5 10 15

Leu Asp Gln Ala Val Gly Glu Ile Asp Glu Ile Glu Ala Gln Arg Ala 20 25 30

Glu Lys Ser Asn Tyr Glu Leu Phe Gln Glu Asp Gly Val Glu Glu His
35 40 45

Thr Arg Pro Ser Tyr Phe Gln Ala Ala Asp Asp Ser Asp Thr Glu Ser 50 55 60

Glu 65	Pro	Glu	Ile	Glu	Asp 70	Asn	Gln	Gly	Leu	Tyr 75	Val	Pro	Asp	Pro	Glu 80
Ala	Glu	Gln	Val	Glu 85	Gly	Phe	Ile	Gln	Gly 90	Pro	Leu	Asp	Asp	Tyr 95	Ala
Asp	Glu	Asp	Val 100	Asp	Val	Val	Phe	Thr 105	Ser	Asp	Trp	Lys	Gln 110	Pro	Glu
Leu	Glu	Ser 115	Asp	Glu	His	Gly	Lys 120	Thr	Leu	Arg	Leu	Thr 125	Leu	Pro	Glu
Gly	Leu 130	Ser	Gly	Glu	Gln	Lys 135	Ser	Gln	Trp	Leu	Leu 140	Thr	Ile	Lys	Ala
Val 145	Val	Gln	Ser	Ala	Lys 150	His	Trp	Asn	Leu	Ala 155	Glu	Cys	Thr	Phe	Glu 160
Ala	Ser	Gly	Glu	Gly 165	Val	Ile	Ile	Lys	Lys 170	Arg	Gln	Ile	Thr	Pro 175	Asp
Val	Tyr	Lys	Val 180	Thr	Pro	Val	Met	Asn 185	Thr	His	Pro	Ser	Gln 190	Ser	Glu
Ala	Val	Ser 195	Asp	Val	Trp	Ser	Leu 200	Ser	Lys	Thr	Ser	Met 205	Thr	Phe	Gln
Pro	Lys 210	Lys	Ala	Ser	Leu	Gln 215	Pro	Leu	Thr	Ile	Ser 220	Leu	Asp	Glu	Leu
Phe 225	Ser	Ser	Arg	Gly	Glu 230	Phe	Ile	Ser	Val	Gly 235	Gly	Asn	Gly	Arg	Met 240
Ser	His	Lys	Glu	Ala 245	Ile	Leu	Leu	Gly	Leu 250	Arg	Tyr	Lys	Lys	Leu 255	Tyr
Asn	Gln	Ala	Arg 260	Val	Lys	Tyr	Ser	Leu 265							
<211)> 27 .> 26 ?> PF	55													
<213	3> V∈	sicu	ılar	ston	natit	is v	rirus	5							
)> 27 Asp		Leu	Thr 5	Lys	Val	Arg	Glu	Туг 10	Leu	Lys	Ser	Tyr	Ser 15	Arg
Leu	Asp	Gln	Ala 20	Val	Gly	Glu	Ile	Asp 25	Glu	Ile	Glu	Ala	Gln 30	Arg	Ala

Glu Lys Ser Asn Tyr Glu Leu Phe Gln Glu Asp Gly Val Glu Glu His 35 40 45

Thr Arg Pro Ser Tyr Phe Gln Ala Ala Asp Asp Ser Asp Thr Glu Ser

50 55 60

Glu Pro Glu Ile Glu Asp Asn Gln Gly Leu Tyr Val Pro Asp Pro Glu 75 70 65 Ala Glu Gln Val Glu Gly Phe Ile Gln Gly Pro Leu Asp Asp Tyr Ala 90 Asp Glu Asp Val Asp Val Val Phe Thr Ser Asp Trp Lys Gln Pro Glu 100 105 Leu Glu Ser Asp Glu His Gly Lys Thr Leu Arg Leu Thr Leu Pro Glu 115 120 Gly Leu Ser Gly Glu Gln Lys Ser Gln Trp Leu Leu Thr Ile Lys Ala Val Val Gln Ser Ala Lys His Trp Asn Leu Ala Glu Cys Thr Phe Glu 145 150 155 160 Ala Ser Gly Glu Gly Val Ile Ile Lys Lys Arg Gln Ile Thr Pro Asp 170 165 Val Tyr Lys Val Thr Pro Val Met Asn Thr His Pro Ser Gln Ser Glu 180 185 Ala Val Ser Asp Val Trp Ser Leu Ser Lys Thr Ser Met Thr Phe Gln 195 200 205 Pro Lys Lys Ala Ser Leu Gln Pro Leu Thr Ile Ser Leu Asp Glu Leu 215 Phe Ser Ser Arg Gly Glu Phe Ile Ser Val Gly Gly Asn Gly Arg Met 230 235 Ser His Lys Glu Ala Ile Leu Leu Gly Leu Arg Tyr Lys Lys Leu Tyr 245 250 Asn Gln Ala Arg Val Lys Tyr Ser Leu 260 265 <210> 28

<211> 690

<212> DNA

<213> Vesicular stomatitis virus

<400> 28

atgagttcct taaagaagat tctcggtctg aaggggaaag gtaagaaatc taagaaatta 60 gggatcgcac cacccctta tgaagaggac actagcatgg agtatgctcc gagcgctcca 120 attgacaaat cctatttgg agttgacgag atggacacct atgatccgaa tcaattaaga 180 tatgagaaat tcttctttac agtgaaaatg acggttagat ctaatcgtcc gttcagaaca 240 tactcagatg tggcagccgc tgtatcccat tgggatcaca tgtacatcgg aatggcaggg 300 aaacgtccct tctacaaaat cttggctttt ttgggttctt ctaatctaaa ggccactcca 360 gcggtattgg cagatcaagg tcaaccagag tatcacactc actgcgaagg cagggcttat 420 ttgccacata ggatgggaa gacccctccc atgctcaatg taccagagca cttcagaaga 480 ccattcaata taggtcttta caagggaacg attgagctca caatgaccat ctacgatgat 540

```
gagtcactgg aagcagctcc tatgatctgg gatcatttca attcttccaa attttctgat 600
ttcagagaga aggccttaat gtttggcctg attgtcgaga aaaaggcatc tggagcgtgg 660
gtcctggatt ctatcagcca cttcaaatga
<210> 29
<211> 690
<212> DNA
<213> Vesicular stomatitis virus
<400> 29
atgagttcct taaagaagat tctcggtctg aaggggaaag gtaagaaatc taagaaatta 60
gggatcgcac cacccctta tgaagaggac actaacatgg agtatgctcc gagcgctcca 120
attgacaaat cctattttgg agttgacgag atggacactc atgatccgca tcaattaaga 180
tatgagaaat tcttctttac agtgaaaatg acggttagat ctaatcgtcc gttcagaaca 240
tactcagatg tggcagccgc tgtatcccat tgggatcaca tgtacatcgg aatggcaggg 300
aaacgtccct tctacaagat cttggctttt ttgggttctt ctaatctaaa ggccactcca 360
gcggtattgg cagatcaagg tcaaccagag tatcacgctc actgtgaagg cagggcttat 420
ttgccacaca gaatggggaa gaccctccc atgctcaatg taccagagca cttcagaaga 480
ccattcaata taggtcttta caagggaacg gttgagctca caatgaccat ctacgatgat 540
gagtcactgg aagcagctcc tatgatctgg gatcatttca attcttccaa attttctgat 600
ttcagagaga aggccttaat gtttggcctg attgtcgaga aaaaggcatc tggagcttgg 660
                                                                  690
gtcctggatt ctgtcagcca cttcaaatga
<210> 30
<211> 690
<212> DNA
<213> Vesicular stomatitis virus
<400> 30
atgagttcct taaagaagat tctcggtctg aaggggaaag gtaagaaatc taagaaatta 60
gggatcgcac caccccctta tgaagaggac actaacatgg agtatgctcc gagcgctcca 120
attgacaaat cctattttgg agttgacgag agggacactc atgatccgca tcaattaaga 180
tatgagaaat tettetttac agtgaaaatg acggttagat etaategtee gtteagaaca 240
tactcagatg tggcagccgc tgtatcccat tgggatcaca tgtacatcgg aatggcaggg 300
aaacgtccct tctacaagat cttggctttt ttgggwtctt ctaatctaaa ggccactcca 360
gcggtattgg cagatcaagg tcaaccagag tatcacgctc actgtgaagg cagggcttat 420
ttgccacaca gaatggggaa gaccctccc atgctcaatg taccagagca cttcagaaga 480
ccattcaata taggtcttta caagggaacg gttgagctca caatgaccat ctacgatgat 540
gagtcactgg aagcagctcc tatgatctgg gatcatttca attcttccaa attttctgat 600
ttcagagaga aggccttaat gtttggcctg attgtcgaga aaaaggcatc tggagcttgg 660
                                                                  690
gtcctggatt ctgtcagcca cttcaaatga
<210> 31
<211> 690
<212> DNA
<213> Vesicular stomatitis virus
<400> 31
atgagttcct taaagaagat tctcggtctg aaggggaaag gtaagaaatc taagaaatta 60
gggatcgcac caccccctta tgaagaggac actaacatgg agtatgctcc gagcgctcca 120
attgacaaat cctattttgg agttgacgag atggacactc atgatccgca tcaattaaga 180
tatgagaaat tottotttac agtgaaaatg acggttagat ctaatcgtcc gttcagaaca 240
tactcagatg tggcagccgc tgtatcccat tgggatcaca tgtacatcgg aatggcaggg 300
aaacgtccct tctacaagat cttggctttt ttgggttctt ctaatctaaa ggccactcca 360
```

geggtattgg cagatcaagg tcaaccagag tatcacgctc actgtgaagg cagggcttat 420 ttgccacaca gaatggggaa gacccctccc atgctcaatg taccagagca cttcagaaga 480 ccattcaata taggtctta caagggaacg gttgagctca caatgaccat ctacgatgat 540 gagtcactgg aagcagctcc tatgatctgg gatcatttca attctccaa attttctgat 600 ttcagagga aggccttaat gtttggcctg attgtcgaga aaaaggcatc tggagcttgg 660 gtcctggatt ctgtcagcca cttcaaatga

<210> 32

<211> 229

<212> PRT

<213> Vesicular stomatitis virus

<400> 32

Met Ser Ser Leu Lys Lys Ile Leu Gly Leu Lys Gly Lys Gly Lys Lys 1 5 10 15

Ser Lys Leu Gly Ile Ala Pro Pro Pro Tyr Glu Glu Asp Thr Ser 20 25 30

Met Glu Tyr Ala Pro Ser Ala Pro Ile Asp Lys Ser Tyr Phe Gly Val 35 40 45

Asp Glu Met Asp Thr Tyr Asp Pro Asn Gln Leu Arg Tyr Glu Lys Phe 50 55 60

Phe Phe Thr Val Lys Met Thr Val Arg Ser Asn Arg Pro Phe Arg Thr 65 70 75 80

Tyr Ser Asp Val Ala Ala Val Ser His Trp Asp His Met Tyr Ile 85 90 95

Gly Met Ala Gly Lys Arg Pro Phe Tyr Lys Ile Leu Ala Phe Leu Gly
100 105 110

Ser Ser Asn Leu Lys Ala Thr Pro Ala Val Leu Ala Asp Gln Gly Gln
115 120 125

Pro Glu Tyr His Thr His Cys Glu Gly Arg Ala Tyr Leu Pro His Arg 130 135 140

Met Gly Lys Thr Pro Pro Met Leu Asn Val Pro Glu His Phe Arg Arg 145 150 155 160

Pro Phe Asn Ile Gly Leu Tyr Lys Gly Thr Ile Glu Leu Thr Met Thr 165 170 175

Ile Tyr Asp Asp Glu Ser Leu Glu Ala Ala Pro Met Ile Trp Asp His 180 185 190

Phe Asn Ser Ser Lys Phe Ser Asp Phe Arg Glu Lys Ala Leu Met Phe 195 200 205

Gly Leu Ile Val Glu Lys Lys Ala Ser Gly Ala Trp Val Leu Asp Ser 210 215 220

Ile Ser His Phe Lys

<210> 33

<211> 229

<212> PRT

<213> Vesicular stomatitis virus

<400> 33

Met Ser Ser Leu Lys Lys Ile Leu Gly Leu Lys Gly Lys Gly Lys Lys

1 5 10 15

Ser Lys Leu Gly Ile Ala Pro Pro Pro Tyr Glu Glu Asp Thr Asn 20 25 30

Met Glu Tyr Ala Pro Ser Ala Pro Ile Asp Lys Ser Tyr Phe Gly Val 35 40 45

Asp Glu Met Asp Thr His Asp Pro His Gln Leu Arg Tyr Glu Lys Phe 50 55 60

Phe Phe Thr Val Lys Met Thr Val Arg Ser Asn Arg Pro Phe Arg Thr 65 70 75 80

Tyr Ser Asp Val Ala Ala Ala Val Ser His Trp Asp His Met Tyr Ile 85 90 95

Gly Met Ala Gly Lys Arg Pro Phe Tyr Lys Ile Leu Ala Phe Leu Gly
100 105 110

Ser Ser Asn Leu Lys Ala Thr Pro Ala Val Leu Ala Asp Gln Gly Gln
115 120 125

Pro Glu Tyr His Ala His Cys Glu Gly Arg Ala Tyr Leu Pro His Arg 130 135 140

Met Gly Lys Thr Pro Pro Met Leu Asn Val Pro Glu His Phe Arg Arg 145 150 155 160

Pro Phe Asn Ile Gly Leu Tyr Lys Gly Thr Val Glu Leu Thr Met Thr 165 170 175

Ile Tyr Asp Asp Glu Ser Leu Glu Ala Ala Pro Met Ile Trp Asp His 180 185 190

Phe Asn Ser Ser Lys Phe Ser Asp Phe Arg Glu Lys Ala Leu Met Phe 195 200 205

Gly Leu Ile Val Glu Lys Lys Ala Ser Gly Ala Trp Val Leu Asp Ser 210 215 220

Val Ser His Phe Lys 225

<210> 34 <211> 229

<212> PRT

<213> Vesicular stomatitis virus

<400> 34

Met Ser Ser Leu Lys Lys Ile Leu Gly Leu Lys Gly Lys Gly Lys Lys 1 5 10 15

Ser Lys Leu Gly Ile Ala Pro Pro Pro Tyr Glu Glu Asp Thr Asn 20 25 30

Met Glu Tyr Ala Pro Ser Ala Pro Ile Asp Lys Ser Tyr Phe Gly Val 35 40 45

Asp Glu Met Asp Thr His Asp Pro His Gln Leu Arg Tyr Glu Lys Phe 50 55 60

Phe Phe Thr Val Lys Met Thr Val Arg Ser Asn Arg Pro Phe Arg Thr 65 70 75 80

Tyr Ser Asp Val Ala Ala Ala Val Ser His Trp Asp His Met Tyr Ile 85 90 95

Gly Met Ala Gly Lys Arg Pro Phe Tyr Lys Ile Leu Ala Phe Leu Gly
100 105 110

Ser Ser Asn Leu Lys Ala Thr Pro Ala Val Leu Ala Asp Gln Gly Gln
115 120 125

Pro Glu Tyr His Ala His Cys Glu Gly Arg Ala Tyr Leu Pro His Arg 130 135 140

Met Gly Lys Thr Pro Pro Met Leu Asn Val Pro Glu His Phe Arg Arg 145 150 155 160

Pro Phe Asn Ile Gly Leu Tyr Lys Gly Thr Val Glu Leu Thr Met Thr 165 170 175

Ile Tyr Asp Asp Glu Ser Leu Glu Ala Ala Pro Met Ile Trp Asp His
180 185 190

Phe Asn Ser Ser Lys Phe Ser Asp Phe Arg Glu Lys Ala Leu Met Phe 195 200 205

Gly Leu Ile Val Glu Lys Lys Ala Ser Gly Ala Trp Val Leu Asp Ser 210 215 220

Val Ser His Phe Lys 225

<210> 35

<211> 229

<212> PRT

<213> Vesicular stomatitis virus

<400> 35

Met Ser Ser Leu Lys Lys Ile Leu Gly Leu Lys Gly Lys Gly Lys Lys

1 5 10 15

Ser Lys Leu Gly Ile Ala Pro Pro Pro Tyr Glu Glu Asp Thr Asn 20 25 30

Met Glu Tyr Ala Pro Ser Ala Pro Ile Asp Lys Ser Tyr Phe Gly Val 35 40 45

Asp Glu Arg Asp Thr His Asp Pro His Gln Leu Arg Tyr Glu Lys Phe
50 55 60

Phe Phe Thr Val Lys Met Thr Val Arg Ser Asn Arg Pro Phe Arg Thr 65 70 75 80

Tyr Ser Asp Val Ala Ala Val Ser His Trp Asp His Met Tyr Ile
85 90 95

Gly Met Ala Gly Lys Arg Pro Phe Tyr Lys Ile Leu Ala Phe Leu Gly
100 105 110

Ser Ser Asn Leu Lys Ala Thr Pro Ala Val Leu Ala Asp Gln Gly Gln 115 120 125

Pro Glu Tyr His Ala His Cys Glu Gly Arg Ala Tyr Leu Pro His Arg 130 135 140

Met Gly Lys Thr Pro Pro Met Leu Asn Val Pro Glu His Phe Arg Arg 145 150 155 160

Pro Phe Asn Ile Gly Leu Tyr Lys Gly Thr Val Glu Leu Thr Met Thr 165 170 175

Ile Tyr Asp Asp Glu Ser Leu Glu Ala Ala Pro Met Ile Trp Asp His
180 185 190

Phe Asn Ser Ser Lys Phe Ser Asp Phe Arg Glu Lys Ala Leu Met Phe 195 200 205

Gly Leu Ile Val Glu Lys Lys Ala Ser Gly Ala Trp Val Leu Asp Ser 210 215 220

Val Ser His Phe Lys 225

<210> 36

<211> 1536

<212> DNA

<213> Vesicular stomatitis virus

<400> 36

atgaagtgcc ttttgtactt agcctttta ttcattgggg tgaattgcaa gttcaccata 60 gtttttcac acaaccaaaa aggaaactgg aaaaatgttc cttctaatta ccattattgc 120 ccgtcaagct cagatttaaa ttggcataat gacttaatag gcacagccat acaagtcaaa 180 atgcccaaga gtcacaaggc tattcaagca gacggttgga tgtgtcatgc ttccaaatgg 240 gtcactactt gtgatttccg ctggtatgga ccgaagtata taacacagtc catccgatcc 300 ttcactccat ctgtagaaca atgcaaggaa agcattgaac aaacgaaaca aggaacttgg 360

```
ctgaatccag gcttccctcc tcaaagttgt ggatatgcaa ctgtgacgga tgccgaagca 420
gtgattgtcc aggtgactcc tcaccatgtg ctggttgatg aatacacagg agaatgggtt 480
gattcacagt tcatcaacgg aaaatgcagc aattacatat gccccactgt ccataactct 540
acaacctggc attctgacta taaggtcaaa gggctatgtg attctaacct catttccatg 600
gacatcacct tcttctcaga ggacggagag ctatcatccc tgggaaagga gggcacaggg 660
ttcagaagta actactttgc ttatgaaact ggaggcaagg cctgcaaaat gcaatactgc 720
aagcattggg gagtcagact cccatcaggt gtctggttcg agatggctga taaggatctc 780
tttgctgcag ccagattccc tgaatgccca gaagggtcaa gtatctctgc tccatctcag 840
acctcagtgg atgtaagtct aattcaggac gttgagagga tcttggatta ttccctctgc 900
caagaaacct ggagcaaaat cagagcgggt cttccaatct ctccagtgga tctcagctat 960
cttgctccta aaaacccagg aaccggtcct gctttcacca taatcaatgg taccctaaaa 1020
tactttgaga ccagatacat cagagtcgat attgctgctc caatcctctc aagaatggtc 1080
ggaatgatca gtggaactac cacagaaagg gaactgtggg atgactgggc accatatgaa 1140
gacgtggaaa ttggacccaa tggagttctg aggaccagtt caggatataa gtttccttta 1200
tacatgattg gacatggtat gttggactcc gatcttcatc ttagctcaaa ggctcaggtg 1260
ttcgaacatc ctcacattca agacgctgct tcgcaacttc ctgatgatga gagtttattt 1320
tttggtgata ctgggctatc caaaaatcca atcgagcttg tagaaggttg gttcagtagt 1380
tggaaaagct ctattgcctc ttttttcttt atcatagggt taatcattgg actattcttg 1440
gttctccgag ttggtatcca tctttgcatt aaattaaagc acaccaagaa aagacagatt 1500
                                                                  1536
tatacagaca tagagatgaa ccgacttgga aagtaa
<210> 37
<211> 885
<212> DNA
<213> Vesicular stomatitis virus
<220>
<221> modified_base
<222> (1)..(885)
<223> "n" represents a, t, c, g, other or unknown
<400> 37
atgaagtgcc ttttgkactt agctttttta ttcatcgggg tgaattgcaa gttcaccata 60
gtttttccat acaaccaaaa aggaaactgg aaaaatgttc cttccaatta ccattattgc 120
ccgtcaagct cagatttaaa ttgncataat gacttaatag gcacagcctt acaagtcaaa 180
atgcccaaga gtcacaaggc tattcaagca gacggttgga tgtgtcatgc ttccaaatgg 240
gtcactactt gtgatttccg ctggtacgga ccgaagtata taacacattc catccgatcc 300
ttcactccat ctgtagaaca atgcaaggaa agcattgaac aaacgaaaca aggaacttgg 360
ctgaatccag gcttccctcc tcaaagttgt ggatatgcaa ctgtgacgga tgctgaagca 420
gcgattgtcc aggtgactcc tcaccatgtg cttgttgatg aatacacagg agaatgggtt 480
gattcacagt tcatcaacgg aaaatgcagc aatgacatat gccccactgt ccataactcc 540
acaacctggc attccgacta taaggtcaaa gggctatgtg attctaacct catttccatg 600
qacatcacct tcttctcaqa qqacqqaqaq ctatcatccc taggaaagga gggcacaggg 660
ttcagaagta actactttgc ttatgaaact ggagacaagg cctgcaaaat gcagtactgc 720
aagcgttggg gagtcagact cccatcaggt gtctggttcg agatggctga taaggatctc 780
tttgctgcag ccagattccc tgaatgccca gaagggtcaa gtatctctgc tccatctcag 840
                                                                  885
acctcagtgg atgtaagtct cattcaggac gttgagagga tcttg
<210> 38
<211> 705
<212> DNA
<213> Vesicular stomatitis virus
<400> 38
ccatctcaga cctcagtgga tgtaagtctc attcaggacg ttgagaggat cttggattat 60
```

```
tccctctgcc aagaaacctg gagcaaaatc agagcgggtc ttcccatctc tccagtggat 120
ctcagctatc ttgctcctaa aaacccagga accggtcctg tctttaccat aatcaatggt 180
accctaaaat actttgagac cagatacatc agagtcgata ttgctgctcc aatcctctca 240
agaatggtcg gaatgatcag tggaactacc acagaaaggg aactgtggga tgactgggct 300
ccatatgaag acgtggaaat tggacccaat ggagttctga ggaccagttc aggatataag 360
tttcctttat atatgattgg acatggtatg ttggactccg atcttcatct tagctcaaag 420
gctcaggtgt ttgaacatcc tcacattcaa gacgctgctt cgcagcttcc tgatgatgag 480
actttatttt ttggtgatac tgggctatcc aaaaatccaa tcgagtttgt agaaggttgg 540
ttcagtagtt ggaagagctc tattgcctct tttttcttta tcatagggtt aatcattgga 600
ctattcttgg ttctccgagt tggtatttat ctttgcatta aattaaagca caccaagaaa 660
                                                                  705
agacagattt atacagacat agagatgaac cgacttggga agtaa
<210> 39
<211> 1536
<212> DNA
<213> Vesicular stomatitis virus
<400> 39
atgaagtgcc ttttgtactt agctttttta ttcatcgggg tgaattgcaa gttcaccata 60
gtttttccat acaaccgaaa aggaaactgg aaaaatgttc cttccaatta ccattattgc 120
ccgtcaagct cagatttaaa ttggcataat gacttaatag gcacagcctt acaagtcaaa 180
atgcccaaga gtcacaaggc tattcaagca gacggttgga tgtgtcatgc ttccaaatgg 240
gtcactactt gtgatttccg ctggtacgga ccgaagtata taacacattc catccgatcc 300
ttcactccat ctgtagaaca atgcaaggaa agcattgaac aaacgaaaca aggaacttgg 360
ctgaatccag gcttccctcc tcaaagttgt ggatatgcaa ctgtgacgga tgctgaagca 420
gcgattgtcc aggtgactcc tcaccatgtg cttgttgatg aatacacagg agaatgggtt 480
gattcacagt tcatcaacgg aaaatgcagc aatgacatat gccccactgt ccataactcc 540
acaacctggc attccgacta taaggtcaaa gggctatgtg attctaacct catttccatg 600
gacatcacct tcttctcaga ggacggagag ctatcatccc taggaaagga gggcacaggg 660
ttcagaagta actactttgc ttatgaaact ggagacaagg cctgcaaaat gcagtactgc 720
aagcattggg gagtcagact cccatcaggt gtctggttcg agatggctga taaggatctc 780
tttgctgcag ccagattccc tgaatgccca gaagggtcaa gtatctctgc tccatctcag 840
acctcagtgg atgtaagtct cattcaggac gttgagagga tcttggatta ttccctctgc 900
caagaaacct ggagcaaaat cagagcgggt cttcccatct ctccagtgga tctcagctat 960
cttgctccta aaaacccagg aaccggtcct gctttcacca taatcaatgg taccctaaaa 1020
tactttgaga ccagatacat cagagtcgat attgctgctc caatcctctc aagaatggtc 1080
ggaatgatca gtggaactac cacagaaagg gaactgtggg atgactgggc tccatatgaa 1140
gacgtggaaa ttggacccaa tggagttctg aggaccagtt caggatataa gtttccttta 1200
tatatgattg gacatggtat gttggactcc gatcttcatc ttagctcaaa ggctcaggtg 1260
tttgaacatc ctcacattca agacgctght gcgcagcttc ctgatgatga gactttattt 1320
tttggtgata ctgggctatc caaaaatcca atcgagtttg tagaaggttg gttcagtagt 1380
tggaagaget ctattgcete ttttttettt atcatagggt taatcattgg actattettg 1440
gttctccgag ttggtattta tctttgcatt aaattaaagc acaccaagaa aagacagatt 1500
                                                                  1536
tatacagaca tagagatgaa ccgacttggg aagtaa
<210> 40
<211> 1399
<212> DNA
<213> Vesicular stomatitis virus
<400> 40
atgaagtgcc ttttgtactt agctttttta ttcatcgggg tgaattgcaa gttcaccata 60
gtttttccat acaaccaaaa aggaaactgg aaaaatgttc cttccaatta ccattattgc 120
ccgtcaagct cagatttaaa ttggcataat gacttaatag gcacagcctt acaggtcaaa 180
atgcccaaga gtcacaaggc tattcaagca gacggttgga tgtgtcatgc ttccaaatgg 240
```

```
gtcactactt gtgatttccg ctggtacgga ccgaagtata taacacattc catccgatcc 300
ttcactccat ctgtagaaca atgcaaggaa agcattgaac aaacgaaaca aggaacttgg 360
ctgaatccag gcttccctcc tcaaagttgt ggatatgcaa ctgtgacgga tgctgaagca 420
gcgattgtcc aggtgactcc tcaccatgtg cttgttgatg aatacacagg agaatgggtt 480
gattcacagt tcatcaacgg aaaatgcagc aatgacatat gccccactgt ccataactcc 540
acaacctggc attccgacta taaggtcaaa gggctatgtg attctaacct catttccatg 600
gacatcacct tetteteaga ggacggagag etateatece taggaaagga gggcacaggg 660
ttcagaagta actactttgc ttatgaaact ggagacaagg cctgcaaaat gcagtactgc 720
aagcgttggg gagtcagact cccatcaggt gtctggttcg ggatggctga taaggatctc 780
tttgctgcag ccagattccc tgaatgccca gaagggtcaa gtatctctgc tccatctcag 840
acctcagtgg atgtaagtct cattcaggac gttgagagga tcttactttg agaccagata 900
catcagagtc gatattgctg ctccaatcct ctcaagaatg gtcggaatga tcagtggaac 960
taccacagaa agggaactgt gggatgactg ggctccatat gaagacgtgg aaattggacc 1020
caatggagtt ctgaggacca gttcaggata taagtttcct ttatatatga ttggacatgg 1080
tatgttggac tccgatcttc atcttagctc aaaggctcag gtgtttgaac atcctcacat 1140
tcaagacgct gcttcgcagc ttcctgatga tgagacttta ttttttggtg atactgggct 1200
atccaaaaat ccaatcgagt ttgtagaagg ttggttcagt agttggaaga gctctattgc 1260
ctcttttttc tttatcatag ggttaatcat tggactattc ttggttctcc gagttggtat 1320
ttatctttgc attaaattaa agcacaccaa gaaaagacag atttatacag acatagagat 1380
                                                                   1399
gaaccgactt gggaagtaa
<210> 41
<211> 511
<212> PRT
<213> Vesicular stomatitis virus
<400> 41
Met Lys Cys Leu Leu Tyr Leu Ala Phe Leu Phe Ile Gly Val Asn Cys
  1
                                     10
Lys Phe Thr Ile Val Phe Pro His Asn Gln Lys Gly Asn Trp Lys Asn
             20
Val Pro Ser Asn Tyr His Tyr Cys Pro Ser Ser Ser Asp Leu Asn Trp
                             40
His Asn Asp Leu Ile Gly Thr Ala Ile Gln Val Lys Met Pro Lys Ser
     50
                         55
His Lys Ala Ile Gln Ala Asp Gly Trp Met Cys His Ala Ser Lys Trp
Val Thr Thr Cys Asp Phe Arg Trp Tyr Gly Pro Lys Tyr Ile Thr Gln
                                                          95
Ser Ile Arg Ser Phe Thr Pro Ser Val Glu Gln Cys Lys Glu Ser Ile
```

105

Glu Gln Thr Lys Gln Gly Thr Trp Leu Asn Pro Gly Phe Pro Pro Gln
115 120 125

Ser Cys Gly Tyr Ala Thr Val Thr Asp Ala Glu Ala Val Ile Val Gln

Val Thr Pro His His Val Leu Val Asp Glu Tyr Thr Gly Glu Trp Val

155

160

135

150

100

130

145

Asp	Ser	Gln	Phe	Ile 165	Asn	Gly	Lys	Cys	Ser 170	Asn	Tyr	Ile	Cys	Pro 175	Thr
Val	His	Asn	Ser 180	Thr	Thr	Trp	His	Ser 185	Asp	Tyr	Lys	Val	Lys 190	Gly	Leu
Cys	Asp	Ser 195	Asn	Leu	Ile	Ser	Met 200	Asp	Ile	Thr	Phe	Phe 205	Ser	Glu	Asp
Gly	Glu 210	Leu	Ser	Ser	Leu	Gly 215	Lys	Glu	Gly	Thr	Gly 220	Phe	Arg	Ser	Asn
Tyr 225	Phe	Ala	Tyr	Glu	Thr 230	Gly	Gly	Lys	Ala	Cys 235	Lys	Met	Gln	Tyr	Cys 240
Lys	His	Trp	Gly	Val 245	Arg	Leu	Pro	Ser	Gly 250	Val	Trp	Phe	Glu	Met 255	Ala
Asp	Lys	Asp	Leu 260	Phe	Ala	Ala	Ala	Arg 265	Phe	Pro	Glu	Cys	Pro 270	Glu	Gly
Ser	Ser	Ile 275	Ser	Ala	Pro	Ser	Gln 280	Thr	Ser	Val	Asp	Val 285	Ser	Leu	Ile
Gln	Asp 290	Val	Glu	Arg	Ile	Leu 295	Asp	Tyr	Ser	Leu	Cys 300	Gln	Glu	Thr	Trp
Ser 305	Lys	Ile	Arg	Ala	Gly 310	Leu	Pro	Ile	Ser	Pro 315	Val	Asp	Leu	Ser	Tyr 320
Leu	Ala	Pro	Lys	Asn 325	Pro	Gly	Thr	Gly	Pro 330	Ala	Phe	Thr	Ile	Ile 335	Asn
Gly	Thr	Leu	Lys 340	Tyr	Phe	Glu	Thr	Arg 345	Tyr	Ile	Arg	Val	Asp 350	Ile	Ala
Ala	Pro	Ile 355	Leu	Ser	Arg	Met	Val 360	Gly	Met	Ile	Ser	Gly 365	Thr	Thr	Thr
Glu	Arg 370	Glu	Leu	Trp	Asp	Asp 375	Trp	Ala	Pro	Tyr	Glu 380	Asp	Val	Glu	Ile
Gly 385	Pro	Asn	Gly	Val	Leu 390	Arg	Thr	Ser	Ser	Gly 395	Tyr	Lys	Phe	Pro	Leu 400
Tyr	Met	Ile	Gly	His 405	Gly	Met	Leu	Asp	Ser 410	Asp	Leu	His	Leu	Ser 415	Ser
Lys	Ala	Gln	Val 420	Phe	Glu	His	Pro	His 425	Ile	Gln	Asp	Ala	Ala 430	Ser	Gln
Leu	Pro	Asp 435	Asp	Glu	Ser	Leu	Phe 440	Phe	Gly	Asp	Thr	Gly 445	Leu	Ser	Lys
Asn	Pro 450	Ile	Glu	Leu	Val	Glu 455	Gly	Trp	Phe	Ser	Ser 460	Trp	Lys	Ser	Ser

Ile Ala Ser Phe Phe Phe Ile Ile Gly Leu Ile Ile Gly Leu Phe Leu 470 Val Leu Arg Val Gly Ile His Leu Cys Ile Lys Leu Lys His Thr Lys 490 Lys Arg Gln Ile Tyr Thr Asp Ile Glu Met Asn Arg Leu Gly Lys 505 500 <210> 42 <211> 295 <212> PRT <213> Vesicular stomatitis virus <220> <221> MOD_RES <222> (1)..(295) <223> "Xaa" represents any, other or unknown amino acid Met Lys Cys Leu Leu Xaa Leu Ala Phe Leu Phe Ile Gly Val Asn Cys 10 Lys Phe Thr Ile Val Phe Pro Tyr Asn Gln Lys Gly Asn Trp Lys Asn Val Pro Ser Asn Tyr His Tyr Cys Pro Ser Ser Ser Asp Leu Asn Xaa 40 His Asn Asp Leu Ile Gly Thr Ala Leu Gln Val Lys Met Pro Lys Ser His Lys Ala Ile Gln Ala Asp Gly Trp Met Cys His Ala Ser Lys Trp 70 Val Thr Thr Cys Asp Phe Arg Trp Tyr Gly Pro Lys Tyr Ile Thr His 85 Ser Ile Arg Ser Phe Thr Pro Ser Val Glu Gln Cys Lys Glu Ser Ile 105 Glu Gln Thr Lys Gln Gly Thr Trp Leu Asn Pro Gly Phe Pro Pro Gln 115 120 Ser Cys Gly Tyr Ala Thr Val Thr Asp Ala Glu Ala Ala Ile Val Gln 135 Val Thr Pro His His Val Leu Val Asp Glu Tyr Thr Gly Glu Trp Val 150 155 Asp Ser Gln Phe Ile Asn Gly Lys Cys Ser Asn Asp Ile Cys Pro Thr 165 170

Val His Asn Ser Thr Thr Trp His Ser Asp Tyr Lys Val Lys Gly Leu

185

190

180

- Cys Asp Ser Asn Leu Ile Ser Met Asp Ile Thr Phe Phe Ser Glu Asp 195 200 205
- Gly Glu Leu Ser Ser Leu Gly Lys Glu Gly Thr Gly Phe Arg Ser Asn 210 215 220
- Tyr Phe Ala Tyr Glu Thr Gly Asp Lys Ala Cys Lys Met Gln Tyr Cys 225 230 235 240
- Lys Arg Trp Gly Val Arg Leu Pro Ser Gly Val Trp Phe Glu Met Ala 245 250 255
- Asp Lys Asp Leu Phe Ala Ala Ala Arg Phe Pro Glu Cys Pro Glu Gly 260 265 270
- Ser Ser Ile Ser Ala Pro Ser Gln Thr Ser Val Asp Val Ser Leu Ile 275 280 285
- Gln Asp Val Glu Arg Ile Leu 290 295
- <210> 43
- <211> 234
- <212> PRT
- <213> Vesicular stomatitis virus
- <400> 43
- Pro Ser Gln Thr Ser Val Asp Val Ser Leu Ile Gln Asp Val Glu Arg
 1 5 10 15
- Ile Leu Asp Tyr Ser Leu Cys Gln Glu Thr Trp Ser Lys Ile Arg Ala
 20 25 30
- Gly Leu Pro Ile Ser Pro Val Asp Leu Ser Tyr Leu Ala Pro Lys Asn 35 40 45
- Pro Gly Thr Gly Pro Val Phe Thr Ile Ile Asn Gly Thr Leu Lys Tyr 50 55 60
- Phe Glu Thr Arg Tyr Ile Arg Val Asp Ile Ala Ala Pro Ile Leu Ser 65 70 75 80
- Arg Met Val Gly Met Ile Ser Gly Thr Thr Thr Glu Arg Glu Leu Trp
 85 90 95
- Asp Asp Trp Ala Pro Tyr Glu Asp Val Glu Ile Gly Pro Asn Gly Val
 100 105 110
- Leu Arg Thr Ser Ser Gly Tyr Lys Phe Pro Leu Tyr Met Ile Gly His
 115 120 125
- Gly Met Leu Asp Ser Asp Leu His Leu Ser Ser Lys Ala Gln Val Phe 130 135 140
- Glu His Pro His Ile Gln Asp Ala Ala Ser Gln Leu Pro Asp Asp Glu

Thr Leu Phe Phe Gly Asp Thr Gly Leu Ser Lys Asn Pro Ile Glu Phe 165 170 175

Val Glu Gly Trp Phe Ser Ser Trp Lys Ser Ser Ile Ala Ser Phe Phe 180 185 190

Phe Ile Ile Gly Leu Ile Ile Gly Leu Phe Leu Val Leu Arg Val Gly 195 200 205

Ile Tyr Leu Cys Ile Lys Leu Lys His Thr Lys Lys Arg Gln Ile Tyr 210 215 220

Thr Asp Ile Glu Met Asn Arg Leu Gly Lys 225 230

<210> 44

<211> 511

<212> PRT

<213> Vesicular stomatitis virus

<220>

<221> MOD_RES

<222> (1)..(511)

<223> "Xaa" represents any, other or unknown amino acid

<400> 44

Met Lys Cys Leu Leu Tyr Leu Ala Phe Leu Phe Ile Gly Val Asn Cys
1 5 10 15

Lys Phe Thr Ile Val Phe Pro Tyr Asn Arg Lys Gly Asn Trp Lys Asn 20 25 30

Val Pro Ser Asn Tyr His Tyr Cys Pro Ser Ser Ser Asp Leu Asn Trp
35 40 45

His Asn Asp Leu Ile Gly Thr Ala Leu Gln Val Lys Met Pro Lys Ser 50 55 60

His Lys Ala Ile Gln Ala Asp Gly Trp Met Cys His Ala Ser Lys Trp 65 70 75 80

Val Thr Thr Cys Asp Phe Arg Trp Tyr Gly Pro Lys Tyr Ile Thr His
85 90 95

Ser Ile Arg Ser Phe Thr Pro Ser Val Glu Gln Cys Lys Glu Ser Ile 100 105 110

Glu Gln Thr Lys Gln Gly Thr Trp Leu Asn Pro Gly Phe Pro Pro Gln
115 120 125

Ser Cys Gly Tyr Ala Thr Val Thr Asp Ala Glu Ala Ala Ile Val Gln 130 135 140

Val Thr Pro His His Val Leu Val Asp Glu Tyr Thr Gly Glu Trp Val

Asp Ser Gln Phe Ile Asn Gly Lys Cys Ser Asn Asp Ile Cys Pro Thr Val His Asn Ser Thr Thr Trp His Ser Asp Tyr Lys Val Lys Gly Leu Cys Asp Ser Asn Leu Ile Ser Met Asp Ile Thr Phe Phe Ser Glu Asp Gly Glu Leu Ser Ser Leu Gly Lys Glu Gly Thr Gly Phe Arg Ser Asn Tyr Phe Ala Tyr Glu Thr Gly Asp Lys Ala Cys Lys Met Gln Tyr Cys Lys His Trp Gly Val Arg Leu Pro Ser Gly Val Trp Phe Glu Met Ala Asp Lys Asp Leu Phe Ala Ala Ala Arg Phe Pro Glu Cys Pro Glu Gly Ser Ser Ile Ser Ala Pro Ser Gln Thr Ser Val Asp Val Ser Leu Ile Gln Asp Val Glu Arg Ile Leu Asp Tyr Ser Leu Cys Gln Glu Thr Trp Ser Lys Ile Arg Ala Gly Leu Pro Ile Ser Pro Val Asp Leu Ser Tyr Leu Ala Pro Lys Asn Pro Gly Thr Gly Pro Ala Phe Thr Ile Ile Asn Gly Thr Leu Lys Tyr Phe Glu Thr Arg Tyr Ile Arg Val Asp Ile Ala Ala Pro Ile Leu Ser Arg Met Val Gly Met Ile Ser Gly Thr Thr Glu Arg Glu Leu Trp Asp Asp Trp Ala Pro Tyr Glu Asp Val Glu Ile Gly Pro Asn Gly Val Leu Arg Thr Ser Ser Gly Tyr Lys Phe Pro Leu Tyr Met Ile Gly His Gly Met Leu Asp Ser Asp Leu His Leu Ser Ser Lys Ala Gln Val Phe Glu His Pro His Ile Gln Asp Ala Xaa Ala Gln Leu Pro Asp Asp Glu Thr Leu Phe Phe Gly Asp Thr Gly Leu Ser Lys

Asn Pro Ile Glu Phe Val Glu Gly Trp Phe Ser Ser Trp Lys Ser Ser

450 455 460

Ile Ala Ser Phe Phe Phe Ile Ile Gly Leu Ile Ile Gly Leu Phe Leu 465 470 475 480

Val Leu Arg Val Gly Ile Tyr Leu Cys Ile Lys Leu Lys His Thr Lys 485 490 495

Lys Arg Gln Ile Tyr Thr Asp Ile Glu Met Asn Arg Leu Gly Lys 500 505 510

<210> 45

<211> 465

<212> PRT

<213> Vesicular stomatitis virus

<400> 45

Met Lys Cys Leu Leu Tyr Leu Ala Phe Leu Phe Ile Gly Val Asn Cys
1 5 10 15

Lys Phe Thr Ile Val Phe Pro Tyr Asn Gln Lys Gly Asn Trp Lys Asn 20 25 30

Val Pro Ser Asn Tyr His Tyr Cys Pro Ser Ser Ser Asp Leu Asn Trp
35 40 45

His Asn Asp Leu Ile Gly Thr Ala Leu Gln Val Lys Met Pro Lys Ser 50 60

His Lys Ala Ile Gln Ala Asp Gly Trp Met Cys His Ala Ser Lys Trp 65 70 75 80

Val Thr Thr Cys Asp Phe Arg Trp Tyr Gly Pro Lys Tyr Ile Thr His
85 90 95

Ser Ile Arg Ser Phe Thr Pro Ser Val Glu Gln Cys Lys Glu Ser Ile 100 105 110

Glu Gln Thr Lys Gln Gly Thr Trp Leu Asn Pro Gly Phe Pro Pro Gln 115 120 125

Ser Cys Gly Tyr Ala Thr Val Thr Asp Ala Glu Ala Ala Ile Val Gln 130 135 140

Val Thr Pro His His Val Leu Val Asp Glu Tyr Thr Gly Glu Trp Val 145 150 155 160

Asp Ser Gln Phe Ile Asn Gly Lys Cys Ser Asn Asp Ile Cys Pro Thr 165 170 175

Val His Asn Ser Thr Thr Trp His Ser Asp Tyr Lys Val Lys Gly Leu 180 185 190

Cys Asp Ser Asn Leu Ile Ser Met Asp Ile Thr Phe Phe Ser Glu Asp 195 200 205

Gly Glu Leu Ser Ser Leu Gly Lys Glu Gly Thr Gly Phe Arg Ser Asn 210 Tyr Phe Ala Tyr Glu Thr Gly Asp Lys Ala Cys Lys Met Gln Tyr Cys 235 Lys Arg Trp Gly Val Arg Leu Pro Ser Gly Val Trp Phe Gly Met Ala 250 Asp Lys Asp Leu Phe Ala Ala Ala Arg Phe Pro Glu Cys Pro Glu Gly 265 260 Ser Ser Ile Ser Ala Pro Ser Gln Thr Ser Val Asp Val Ser Leu Ile 280 Gln Asp Val Glu Arg Ile Tyr Phe Glu Thr Arg Tyr Ile Arg Val Asp 290 295 Ile Ala Ala Pro Ile Leu Ser Arg Met Val Gly Met Ile Ser Gly Thr Thr Thr Glu Arg Glu Leu Trp Asp Asp Trp Ala Pro Tyr Glu Asp Val 330 Glu Ile Gly Pro Asn Gly Val Leu Arg Thr Ser Ser Gly Tyr Lys Phe 340 345 Pro Leu Tyr Met Ile Gly His Gly Met Leu Asp Ser Asp Leu His Leu 360 Ser Ser Lys Ala Gln Val Phe Glu His Pro His Ile Gln Asp Ala Ala 375 370 Ser Gln Leu Pro Asp Asp Glu Thr Leu Phe Phe Gly Asp Thr Gly Leu 390 Ser Lys Asn Pro Ile Glu Phe Val Glu Gly Trp Phe Ser Ser Trp Lys 410 Ser Ser Ile Ala Ser Phe Phe Phe Ile Ile Gly Leu Ile Ile Gly Leu 420 425 430 Phe Leu Val Leu Arg Val Gly Ile Tyr Leu Cys Ile Lys Leu Lys His 440 Thr Lys Lys Arg Gln Ile Tyr Thr Asp Ile Glu Met Asn Arg Leu Gly 450 455 460

Lys 465

<210> 46

<211> 6330

<212> DNA

<213> Vesicular stomatitis virus

atggaagtcc acgattttga gaccgacgag ttcaatgatt tcaatgaaga tgactatgcc 60 acaagagaat teetgaatee egatgagege atgaegtaet tgaateatge tgattacaat 120 ttgaattctc ctctaattag tgatgatatt gacaatttga tcaggaaatt caattctctt 180 ccgattccct cgatgtggga tagtaagaac tgggatggag ttcttgagat gttaacatca 240 tqtcaaqcca atcccatctc aacatctcag atgcataaat ggatgggaag ttggttaatg 300 tctgataatc atgatgccag tcaagggtat agttttttac atgaagtgga caaagaggca 360 gaaataacat ttgacgtggt ggagaccttc atccgcggct ggggcaacaa accaattgaa 420 tacatcaaaa aggaaagatg gactgactca ttcaaaattc tcgcttattt gtgtcaaaag 480 tttttqqact tacacaagtt gacattaatc ttaaatgctg tctctgaggt ggaattgctc 540 aacttggcga ggactttcaa aggcaaagtc agaagaagtt ctcatggaac gaacatatgc 600 aggattaggg ttcccagctt gggtcctact tttatttcag aaggatgggc ttacttcaag 660 aaacttgata ttctaatgga ccgaaacttt ctgttaatgg tcaaagatgt gattataggg 720 aggatgcaaa cggtgctatc catggtatgt agaatagaca acctgttctc agagcaagac 780 atcttctccc ttctaaatat ctacagaatt ggagataaaa ttgtggagag gcagggaaat 840 ttttcttatg acttgattaa aatggtggaa ccgatatgca acttgaagct gatgaaatta 900 gcaagagaat caaggccttt agtcccacaa ttccctcatt ttgaaaatca tatcaagact 960 tctgttgatg aaggggcaaa aattgaccga ggtataagat tcctccatga tcagataatg 1020 agtgtgaaaa cagtggatct cacactggtg atttatggat cgttcagaca ttggggtcat 1080 ccttttatag attattacac tggactagaa aaattacatt cccaagtaac catgaagaaa 1140 gatattgatg tgtcatatgc aaaagcactt gcaagtgatt tagctcggat tgttctattt 1200 caacagttca atgatcataa aaagtggttc gtgaatggag acttgctccc tcatgatcat 1260 ccctttaaaa gtcatgttaa agaaaataca tggcccacag ctgctcaagt tcaagatttt 1320 ggagataaat ggcatgaact tccgctgatt aaatgttttg aaatacccga cttactagac 1380 ccatcgataa tatactctga caaaagtcat tcaatgaata ggtcagaggt gttgaaacat 1440 gtccgaatga atccgaacac tcctatccct agtaaaaagg tgttgcagac tatgttggac 1500 acaaaggcta ccaattggaa agaatttctt aaagagattg atgagaaggg cttagatgat 1560 gatgatctaa ttattggtct taaaggaaag gagagggaac tgaagttggc aggtagattt 1620 ttctccctaa tgtcttggaa attgcgagaa tactttgtaa ttaccgaata tttgataaag 1680 actcatttcg tccctatgtt taaaggcctg acaatggcgg acgatctaac tgcagtcatt 1740 aaaaagatgt tagattcctc atccggccaa ggattgaagt catatgaggc aatttgcata 1800 gccaatcaca ttgattacga aaaatggaat aaccaccaaa ggaagttatc aaacggccca 1860 gtgttccgag ttatgggcca gttcttaggt tatccatcct taatcgagag aactcatgaa 1920 ttttttgaga aaagtcttat atactacaat ggaagaccag acttgatgcg tgttcacaac 1980 aacacactga tcaattcaac ctcccaacga gtttgttggc aaggacaaga gggtggactg 2040 gaaggtctac ggcaaaaagg atggactatc ctcaatctac tggttattca aagagaggct 2100 aaaatcagaa acactgctgt caaagtcttg gcacaaggtg ataatcaagt tatttgcaca 2160 cagtataaaa cgaagaaatc gagaaacgtt gtagaattac agggtgctct caatcaaatg 2220 gtttctaata atgagaaaat tatgactgca atcaaaatag ggacagggaa gttaggactt 2280 ttgataaatg acgatgagac tatgcaatct gcagattact tgaattatgg aaaaataccg 2340 attttccgtg gagtgattag agggttagag accaagagat ggtcacgagt gacttgtgtc 2400 accaatgacc aaatacccac ttgtgctaat ataatgagct cagtttccac aaatgctctc 2460 accgtagete attttgetga gaacceaate aatgceatga tacagtacaa ttattttggg 2520 acatttgcta gactcttgtt gatgatgcat gatcctgctc ttcgtcaatc attgtatgaa 2580 gttcaagata agataccggg cttgcacagt tctactttca aatacgccat gttgtatttg 2640 gaccetteca ttggaggagt gtegggeatg tetttgteca ggtttttgat tagageette 2700 ccagatcccg taacagaaag tctctcattc tggagattca tccatgtaca tgctcgaagt 2760 gagcatctga aggagatgag tgcagtattt ggaaaccccg agatagccaa gtttcgaata 2820 actcacatag acaagctagt agaagatcca acctctctga acatcgctat gggaatgagt 2880 ccagcgaact tgttaaagac tgaggttaaa aaatgcttaa tcgaatcaag acaaaccatc 2940 aggaaccagg tgattaagga tgcaaccata tatttgtatc atgaagagga tcggctcaga 3000 agtttcttat ggtcaataaa tcctctgttc cctagatttt taagtgaatt caaatcaggc 3060 actititing gagtegeaga egggeteate agtetatite aaaatteteg tactattegg 3120 aactccttta agaaaaagta tcatagggaa ttggatgatt tgattgtgag gagtgaggta 3180 tcctctttga cacatttagg gaaacttcat ttgagaaggg gatcatgtaa aatgtggaca 3240 tgttcagcta ctcatgctga cacattaaga tacaaatcct ggggccgtac agttattggg 3300 acaactgtac cccatccatt agaaatgttg ggtccacaac atcgaaaaga gactccttgt 3360

```
gcaccatgta acacatcagg gttcaattat gtttctgtgc attgtccaga cgggatccat 3420
gacgtcttta gttcacgggg accattgcct gcttatctag ggtctaaaac atctgaatct 3480
acatctattt tgcagccttg ggaaagggaa agcaaagtcc cactgattaa aagagctaca 3540
cgtcttagag atgctatctc ttggtttgtt gaacccgact ctaaactagc aatgactata 3600
ctttctaaca tccactcttt aacaggcgaa gaatggacca aaaggcagca tgggttcaaa 3660
agaacagggt ctgcccttca taggttttcg acatctcgga tgagccatgg tgggttcgca 3720
tctcagagca ctgcagcatt gaccaggttg atggcaacta cagacaccat gagggatctg 3780
ggagatcaga atttcgactt tttattccaa gcaacgttgc tctatgctca aattaccacc 3840
actgttgcaa gagacggatg gatcaccagt tgtacagatc attatcatat tgcctgtaag 3900
tectgtttga gacccataga agagateace etggaeteaa gtatggaeta eaegeeecea 3960
gatgtatccc atgtgctgaa gacatggagg aatggggaag gttcgtgggg acaagagata 4020
aaacagatct atcctttaga agggaattgg aagaatttag cacctgctga gcaatcctat 4080
caagtcggca gatgtatagg ttttctatat ggagacttgg cgtatagaaa atctactcat 4140
gccgaggaca gttctctatt tcctctatct atacaaggtc gtattagagg tcgaggtttc 4200
ttaaaagggt tgctagacgg attaatgaga gcaagttgct gccaagtaat acaccggaga 4260
agtetggete atttgaagag geeggeeaac geagtgtaeg gaggtttgat ttaettgatt 4320
gataaattga gtgtatcacc tccattcctt tctcttacta gatcaggacc tattagagac 4380
gaattagaaa cgattcccca caagatccca acctcctatc cgacaagcaa ccgtgatatg 4440
ggggtgattg tcagaaatta cttcaaatac caatgccgtc taattgaaaa gggaaaatac 4500
agatcacatt attcacaatt atggttattc tcagatgtct tatccataga cttcattgga 4560
ccattctcta tttccaccac cctcttgcaa atcctataca agccattttt atctgggaaa 4620
gataagaatg agttgagaga gctggcaaat ctttcttcat tgctaagatc aggagagggg 4680
tgggaagaca tacatgtgaa attcttcacc aaggacatat tattgtgtcc agaggaaatc 4740
agacatgctt gcaagttcgg gattgctaag gataataata aagacatgag ctatcccct 4800
tggggaaggg aatccagagg gacaattaca acaatccctg tttattatac gaccacccct 4860
tacccaaaga tgctagagat gcctccaaga atccaaaatc ccctgctgtc cggaatcagg 4920
ttgggccaat taccaactgg cgctcattat aaaattcgga gtatattaca tggaatggga 4980
atccattaca gggacttctt gagttgtgga gacggctccg gagggatgac tgctgcatta 5040
ctacgagaaa atgtgcatag cagaggaata ttcaatagtc tgttagaatt atcagggtca 5100
gtcatgcgag gcgcctctcc tgagcccccc agtgccctag aaactttagg aggagataaa 5160
togagatgtg taaatggtga aacatgttgg gaatatccat ctgacttatg tgacccaagg 5220
acttgggact atttcctccg actcaaagca ggcttggggc ttcaaattga tttaattgta 5280
atggatatgg aagtteggga ttettetaet ageetgaaaa ttgagaegaa tgttagaaat 5340
tatgtgcacc ggattttgga tgagcaagga gttttaatct acaagactta tggaacatat 5400
atttgtgaga gcgaaaagaa tgcagtaaca atccttggtc ccatgttcaa gacggtcgac 5460
ttagttcaaa cagaatttag tagttctcaa acgtctgaag tatatatggt atgtaaaggt 5520
ttgaagaaat taatcgatga acccaatccc gattggtctt ccatcaatga atcctggaaa 5580
aacctgtacg cattccagtc atcagaacag gaatttgcca gagcaaagaa ggttagtaca 5640
tactttacct tgacaggtat tccctcccaa ttcattcctg atccttttgt aaacattgag 5700
actatgetac aaatattegg agtaceeacg ggtgtgtete atgeggetge ettaaaatea 5760
tctgatagac ctgcagattt attgaccatt agcctttttt atatggcgat tatatcgtat 5820
tataacatca atcatatcag agtaggaccg atacctccga accccccatc agatggaatt 5880
gcacaaaatg tggggatcgc tataactggt ataagctttt ggctgagttt gatggagaaa 5940
gacattccac tatatcaaca gtgtttagca gttatccagc aatcattccc gattaggtgg 6000
gaggctgttt cagtaaaagg aggatacaag cagaagtgga gtactagagg tgatgggctc 6060
ccaaaagata cccgaacttc agactccttg gccccaatcg ggaactggat cagatctctg 6120
gaattggtcc gaaaccaagt tcgtctaaat ccattcaatg agatcttgtt caatcagcta 6180
tgtcgtacag tggataatca tttgaaatgg tcaaatttgc gaagaaacac aggaatgatt 6240
gaatggatca atagacgaat ttcaaaagaa gaccggtcta tactgatgtt gaagagtgac 6300
                                                                  6330
ctacacgagg aaaactcttg gagagattaa
```

<210> 47 <211> 5327

<212> DNA

<213> Vesicular stomatitis virus

<400> 4/						
cattggggtc	atccttttat	agattattac	gctggwctag	aaaaattaca	ttcccaagtw	60
accatkaaga	aagatattga	tgtgtcatat	gcraaagcac	ttgcaagtga	tttagctcgg	120
attgttctat	ttcaacagtt	caatgatcat	amaaagtggt	tcgtgaatgg	agacttgctc	180
cctcatgatc	atccctttaa	aagtcatgtt	aaagaaaata	catggcccac	agctgctcaa	240
gttcaagatt	ttggagataa	atggcatgaa	cttccgctga	ttaaatgttt	tgaaataccc	300
		aatatactct				
		gaatccgaac				
		taccaattgg				
		aattattggt				
		aatgtcttgg				
		cgtccctatg				
		gttagattcc				
		cattgattac				
		agttatgggc				
		gaaaagtctt				
		gatcaattca				
		acggcaaaaa				
		aaacactgct				
		aacgaagaaa				
		taatgagaaa				
		tgacgatgag				
		tggagtgatt				
		ccaaataccc				
		tcattttgct				
		tagactcttg				
		taagataccg				
		cattggagga				
		cgtaacagaa				
		gaaggagatg				
		agacaagcta				
		cttgttaaag				
		ggtgattaag				
		atggtcaata				
		gggagtcgca				
		taagaaaaag				
aggagtgagg	tatcctcttt	gacacattta	gggaaacttc	atttgagaag	gggatcatgt	2160
aaaatgtgga	catgttcagc	tactcatgct	gacacattaa	gatacaaatc	ctggggccgt	2220
		accccatcca				
		taacacatca				
		tagttcacgg				
		tttgcagcct				
		agatgctatc				
		catccactct				
		gtctgccctt				
		cactgcagca				
		gaatttcgac				
		aagagacgga				
attacctata	agtcctgttt	gagacccata	gaagagatca	ccctggactc	aagtatggac	2880
		ccatgtgctg				
		ctatccttta				
		cagatgtata				
		cagttctcta				
		gttgctagac				
		tcatttgaag				
		gagtgtatca				
cctattagag	acyaattaga	aacgattccc	cacaagaccc	Caaceteeta	cccyacaage	2200

```
aaccgtgata tgggggtgat tgtcagaaat tacttcaaat accaatgccg tctaattgaa 3420
aagggaaaat acagatcaca ttattcacaa ttatggttat tctcagatgt cttatccata 3480
qacttcattg gaccattctc tatttccacc accetcttgc aaatcctata caagccattt 3540
ttatctggga aagataagaa tgagttgaga gagctggcaa atctttcttc attgctaaga 3600
tcaggagagg ggtgggaaga catacatgtg aaattcttca ccaaggacat attattgtgt 3660
ccaqaqqaaa tcagacatgc ttgcaagttc gggattgcta aggataataa taaagacatg 3720
agctatcccc cttggggaag ggaatccaga gggacaatta caacaatccc tgtttattat 3780
acgaccaccc cttacccaaa gatgctagag atgcctccaa gaatccaaaa tcccctgctg 3840
tccggaatca ggttgggcca gttaccaact ggcgctcatt ataaaattcg gagtatatta 3900
catggaatgg gaatccatta cagggacttc ttgagttgtg gagacggctc cggagggatg 3960
actgctgcat tactacgaga aaatgtgcat agcagaggaa tattcaatag tctgttagaa 4020
ttatcagggt cagtcatgcg aggcgcctct cctgagcccc ccagtgccct agaaacttta 4080
ggaggagata aatcgagatg tgtaaatggt gaaacatgtt gggaatatcc atctgactta 4140
tgtgacccaa ggacttggga ctatttcctc cgactcaaag caggcttggg gcttcaaatt 4200
gatttaattg taatggatat ggaagttcgg gattcttcta ctagcctgaa aattgagacg 4260
aatgttagaa attatgtgca ccggattttg gatgagcaag gagttttaat ctacaagact 4320
tatggaacat atatttgtga gagcgaaaag aatgcagtaa caatccttgg tcccatgttc 4380
aagacggtcg acttagttca aacagaattt agtagttctc aaacgtctga agtatatatg 4440
gtatgtaaag gtttgaagaa attaatcgat gaacccaatc ccgattggtc ttccatcaat 4500
gaatcctgga aaaacctgta cgcattccag tcatcagaac aggaatttgc cagagcaaag 4560
aaggttagta catactttac cttgacaggt attccctccc aattcattcc tgatcctttt 4620
gtgaacattg agactatgct acaaatattc ggagtaccca cgggtgtgtc tcatgcggct 4680
gccttaaaat catctgatag acctgcagat ttattgacca ttagcctttt ttatatggcg 4740
attatatcqt attataacat caatcatatc agagtaggac cgatacctcc gaacccccca 4800
tcagatggaa ttgcacaaaa tgtggggatc gctataactg gtataagctt ttggctgagt 4860
ttgatggaga aagacattcc actatatcaa cagtgtttag cagttatcca gcaatcattc 4920
ccgattaggt gggaggctgt ttcagtaaaa ggaggataca agcagaagtg gagtactaga 4980
ggtgatgggc tcccaaaaga tacccgaatt tcagactcct tggccccaat cgggaactgg 5040
atcagatete tqqaattqqt eeqaaaccaa qtteqtetaa atceatteaa tgagatettg 5100
ttcaatcagc tatgtcgtac agtggataat catttgaaat ggtcaaattt gcgaaaaaac 5160
acaggaatga ttgaatggat caatagacga atttcaaaag aagaccggtc tatactgatg 5220
ttgaagagtg acctacatga ggaaaactct tggagagatt aaaaaatcat gaggagactc 5280
caaactttaa gtatgaaaaa aactttgatc cttaagaccc tcttgtg
                                                                  5327
<210> 48
<211> 348
<212> DNA
<213> Vesicular stomatitis virus
<400> 48
atggaagtcc acgattttga gaccgacgag ttcaatgatt tcaatgaaga tgactatgcc 60
acaagagaat teetgaatee egatgagege atgaegtaet tgaateatge tgattacaat 120
ttgaattctc ctctaattag tgatgatatt gacaatttga tcaggaaatt caattctctt 180
ccgattccct cgatgtggga tagtaagaac tgggatggag ttcttgagat gttaacatca 240
tgtcaagcca atcccatctc aacatctcag atgcataaat ggatgggaag ttggttaatg 300
                                                                  348
tctgataatc atgatgccag tcaagggtat agttttttac atgaagtg
<210> 49
<211> 6330
```

<220>
<221> modified_base
<222> (1)..(6330)

<213> Vesicular stomatitis virus

<212> DNA

<400> 49 atggaagtcc acgattttga gaccgacgag ttcaatgatt tcaatgaaga tgactatgcc 60 acaagagaat teetgaatee egatgagege atgaegtaet tgaateatge tgattacaat 120 ttgaattctc ctctaattag tgatgatatt gacaatttga tcaggaaatt caattctctt 180 ccgattccct cgatgtggga tagtaagaac tgggatggag ttcttgagat gttaacatca 240 tgtcaagcca atcccatctc aacatctcag atgcataaat ggatgggaag ttggttaatg 300 tctgataatc atgatgccag tcaagggtat agttttttac atgaagtgga caaagaggca 360 gaaataacat ttgacgtggt ggagaccttc atccgcggct ggggcaacaa accaattgaa 420 tacatcaaaa aggaaagatg gactgactca ttcaaaattc tcgcttattt gtgtcaaaag 480 ttttttggact tacacaagtt gacattaatc ttaaatgctg tctctgaggt ggaattgctc 540 aacttggcga ggactttcaa aggcaaagtc agaagaagtt ctcatggaac gaacatatgc 600 aggettaggg tteccagett gggteetaet tttattteag aaggatggge ttaetteaag 660 aaacttgata ttctaatgga ccgaaacttt ctgttaatgg tcaaagatgt gattataggg 720 aggatgcaaa cggtgctatc catggtatgt agaatagaca acctgttctc agagcaagac 780 atcttctccc ttctaaatat ctacagaatt ggagataaaa ttgtggagag gcagggaaat 840 ttttcttatg acttgattaa aatggtggaa ccgatatgca acttgaggct gatgaaatta 900 gcaagagaat caaggcettt agteecacaa tteecteatt ttgaaaatea tateaagaet 960 tctgttgatg aaggggcaaa aattgaccga ggtataagat tcctccatga tcagataatg 1020 agtgtgaaaa cagtggatct cacactggtg atttatggat cgttcagaca ttggggtcat 1080 ccttttatag attattacgc tggactagaa aaattacatt cccaagtaac catgaagaaa 1140 gatattgatg tgtcatatgc aaaagcactt gcaagtgatt tagctcggat tgttctattt 1200 caacagttca atgatcataa aaagtggttc gtgaatggag acttgctccc tcatgatcat 1260 ccctttaaaa gtcatgttaa agaaaataca tggcccacag ctgctcaagt tcaagatttt 1320 ggagataaat ggcatgaact tccgctgatt aaatgttttg aaatacccga cttactagac 1380 ccatcgataa tatactctga caaaagtcat tcaatgaata ggtcagaggt gttgaaacat 1440 gtccgaatga atccgaacac tcctatccct agtaaaaaagg tgttgcagac tatgttggac 1500 acaaaggcta ccaattggaa agaatttctt aaagagattg atgagaaggg cttagatgat 1560 gatgatctaa ttattggtct taaaggaaag gagagggaac tgaagttggc aggtagattt 1620 ttctccctaa tgtcttggaa attgcgagaa tactttgtaa ttaccgaata tttgataaag 1680 actcatttcg tccctatgtt taaaggcctg acaatggcgg acgatctaac tgcagtcatt 1740 aaaaagatgt tagattcctc atccggccaa ggattgaagt catatgaggc aatttgcata 1800 gccaatcaca ttgattacga aaaatggaat aaccaccaaa ggaagttatc aaacggccca 1860 gtgttccgag ttatgggcca gttcttaggt tatccatcct taatcgagag aactcatgaa 1920 tttttttgaga aaagtettat atactacaat ggaagaccag acttgatgeg tgttcacaac 1980 aacacactga tcaattcaac ctcccaacga gtttgttggc aaggacaaga gggtggactg 2040 gaaggtctac ggcaaaaagg atggagtatc ctcaatctac tggttattca aagagaggct 2100 aaaatcagaa acactgctgt caaagtcttg gcacaaggtg ataatcaagt tatttgcaca 2160 cagtataaaa cgaagaaatc gagaaacgtt gtagaattac agggtgctct caatcaaatg 2220 gtttctaata atgagaaaat tatgactgca atcaaaatag ggacagggaa gttaggactt 2280 ttgataaatg acgatgagac tatgcaatct gcagattact tgaattatgg aaaaatacca 2340 attttccgtg gagtgattag agggttagag accaagagat qqtcacgagt gacttgtgtc 2400 accaatgacc aaatacccac ttgtgctaat ataatgagct caqtttccac aaatgctctc 2460 accgtagctc attttgctga gaacccaatc aatgccatga tacagtacaa ttattttggg 2520 acatttgcta gactcttgtt gatgatgcat gatcctgctc ttcgtcaatc attgtatgaa 2580 gttcaagata agataccggg cttgcacagt tctactttca aatacgccat gttgtatttg 2640 gaccetteca ttggaggagt gtegggeatg tetttgteea ggtttttgat tagageette 2700 ccagatcccg taacagaaag tctctcattc tggagattca tccatgtaca tgctcgaagt 2760 gagcatctga aggagatgag tgcagtattt ggaaaccccg agatagccaa gtttcgaata 2820 actcacatag acaagctagt agaagatcca acctctctga acatcgctat gggaatgagt 2880 ccagcgaact tgttaaagac tgaggttaaa aaatgcttaa tcgaatcaag acaaaccatc 2940 aggaaccagg tgattaagga tgcaaccata tatttgtatc atgaagagga tcggctcaga 3000 agtttcttat ggtcaataaa tcctctgttc cctagatttt taagtgaatt caaatcaggc 3060

acttttttgg gagtcgcaga cgggctcatc agtctatttc aaaattctcg tactattcgg 3120 aactccttta agaaaaagta tcatagggaa ttggatgatt tgattgtgag gagtgaggta 3180 tcctctttga cacatttagg gaaacttcat ttgagaaggg gatcatgtaa aatgtggaca 3240

tgttcagcta	ctcatgctga	cacattaaga	tacaaatcct	ggggccgtac	agttattggg	3300
acaactgtac	cccatccatt	agaaatgttg	ggtccacaac	atcgaaaaga	gactccttgt	3360
gcaccatgta	acacatcagg	gttcaattat	gtttctgtgc	attgtccaga	cgggatccat	3420
gacgtcttta	gttcacgggg	accattgcct	gcttatctag	ggtctaaaac	atctgaatct	3480
		ggaaagggaa				
		ttggtttgtt				
		aacaggcgaa				
		taggttttcg				
		gaccaggttg				
		tttattccan				
		gatcaccagt				
		agagatcacc				
gatgtatccc	atgtgctgaa	gacatggagg	aatggggaag	gttcgtgggg	acaagagata	4020
		agggaattgg				
		ttttctatat				
		tcctctatct				
		attaatgaga				
		gccggccaac				
gataaattga	gtgtatcacc	tccattcctt	tctcttacta	gatcaggacc	tattagagac	4380
		caagatccca				
		cttcaaatac				
agatcacatt	attcacaatt	atggttattc	tcagatgtct	tatccataga	cttcattgga	4560
ccattctcta	tttccaccac	cctcttgcaa	atcctataca	agccattttt	atctgggaaa	4620
gataagaatg	agttgagaga	gctggcaaat	ctttcttcat	tgctaagatc	aggagagggg	4680
tgggaagaca	tacatgtgaa	attcttcacc	aaggacatat	tattgtgtcc	agaggaaatc	4740
agacatgctt	gcaagttcgg	gattgctaag	gataataata	aagacatgag	ctatccccct	4800
tggggaaggg	aatccagagg	gacaattaca	acaatccctg	tttattatac	gaccacccct	4860
tacccaaaga	tgctagagat	gcctccaaga	atccaaaatc	ccctgctgtc	cggaatcagg	4920
ttgggccagt	taccaactgg	cgctcattat	aaaattcgga	gtatattaca	tggaatggga	4980
atccattaca	gggacttctt	gagttgtgga	gacggctccg	gagggatgac	tgctgcatta	5040
ctacgagaaa	atgtgcatag	cagaggaata	ttcaatagtc	tgttagaatt	atcagggtca	5100
		tgagcccccc				
tcgagatgtg	taaatggtga	aacatgttgg	gaatatccat	ctgacttatg	tgacccaagg	5220
		actcaaagca				
		ttcttctact				
tatgtgcacc	ggattttgga	tgagcaagga	gttttaatct	acaagactta	tggaacatat	5400
		tgcagtaaca				
		tagttctcaa				
		acccaatccc				
		atcagaacag				
		tccctcccaa				
		agtacccacg				
		attgaccatt				
		agtaggaccg				
		tataactggt				
		gtgtttagca				
		aggatacaag				
		agactccttg				
		tcgtctaaat				
		tttgaaatgg				
		ttcaaaagaa				
	aaaactcttg			5 5 -		6330
		J J J				•

<400> 50

Met Glu Val His Asp Phe Glu Thr Asp Glu Phe Asn Asp Phe Asn Glu
1 5 10 15

Asp Asp Tyr Ala Thr Arg Glu Phe Leu Asn Pro Asp Glu Arg Met Thr 20 25 30

Tyr Leu Asn His Ala Asp Tyr Asn Leu Asn Ser Pro Leu Ile Ser Asp 35 40 45

Asp Ile Asp Asn Leu Ile Arg Lys Phe Asn Ser Leu Pro Ile Pro Ser 50 55 60

Met Trp Asp Ser Lys Asn Trp Asp Gly Val Leu Glu Met Leu Thr Ser 65 70 75 80

Cys Gln Ala Asn Pro Ile Ser Thr Ser Gln Met His Lys Trp Met Gly
85 90 95

Ser Trp Leu Met Ser Asp Asn His Asp Ala Ser Gln Gly Tyr Ser Phe 100 105 110

Leu His Glu Val Asp Lys Glu Ala Glu Ile Thr Phe Asp Val Val Glu
115 120 125

Thr Phe Ile Arg Gly Trp Gly Asn Lys Pro Ile Glu Tyr Ile Lys Lys 130 135 140

Glu Arg Trp Thr Asp Ser Phe Lys Ile Leu Ala Tyr Leu Cys Gln Lys 145 150 155 160

Phe Leu Asp Leu His Lys Leu Thr Leu Ile Leu Asn Ala Val Ser Glu 165 170 175

Val Glu Leu Leu Asn Leu Ala Arg Thr Phe Lys Gly Lys Val Arg Arg 180 185 190

Ser Ser His Gly Thr Asn Ile Cys Arg Ile Arg Val Pro Ser Leu Gly 195 200 205

Pro Thr Phe Ile Ser Glu Gly Trp Ala Tyr Phe Lys Lys Leu Asp Ile 210 215 220

Leu Met Asp Arg Asn Phe Leu Leu Met Val Lys Asp Val Ile Ile Gly 225 230 235 240

Arg Met Gln Thr Val Leu Ser Met Val Cys Arg Ile Asp Asn Leu Phe 245 250 255

Ser Glu Gln Asp Ile Phe Ser Leu Leu Asn Ile Tyr Arg Ile Gly Asp 260 265 270

Lys Ile Val Glu Arg Gln Gly Asn Phe Ser Tyr Asp Leu Ile Lys Met 275 280 285

Val	Glu 290	Pro	Ile	Cys	Asn	Leu 295	Lys	Leu	Met	Lys	Leu 300	Ala	Arg	Glu	Ser
Arg 305	Pro	Leu	Val	Pro	Gln 310	Phe	Pro	His	Phe	Glu 315	Asn	His	Ile	Lys	Thr 320
Ser	Val	Asp	Glu	Gly 325	Ala	Lys	Ile	Asp	Arg 330	Gly	Ile	Arg	Phe	Leu 335	His
Asp	Gln	Ile	Met 340	Ser	Val	Lys	Thr	Val 345	Asp	Leu	Thr	Leu	Val 350	Ile	Tyr
Gly	Ser	Phe 355	Arg	His	Trp	Gly	His 360	Pro	Phe	Ile	Asp	Tyr 365	Tyr	Thr	Gly
Leu	Glu 370	Lys	Leu	His	Ser	Gln 375	Val	Thr	Met	Lys	Lys 380	Asp	Ile	Asp	Val
Ser 385	Tyr	Ala	Lys	Ala	Leu 390	Ala	Ser	Asp	Leu	Ala 395	Arg	Ile	Val	Leu	Phe 400
Gln	Gln	Phe	Asn	Asp 405	His	Lys	Lys	Trp	Phe 410	Val	Asn	Gly	Asp	Leu 415	Leu
Pro	His	Asp	His 420	Pro	Phe	Lys	Ser	His 425	Val	Lys	Glu	Asn	Thr 430	Trp	Pro
Thr	Ala	Ala 435	Gln	Val	Gln	Asp	Phe 440	Gly	Asp	Lys	Trp	His 445	Glu	Leu	Pro
Leu	Ile 450	Lys	Cys	Phe	Glu	Ile 455	Pro	Asp	Leu	Leu	Asp 460	Pro	Ser	Ile	Ile
Туг 465	Ser	Asp	Lys	Ser	His 470	Ser	Met	Asn	Arg	Ser 475	Glu	Val	Leu	Lys	His 480
Val	Arg	Met	Asn	Pro 485	Asn	Thr	Pro	Ile	Pro 490	Ser	Lys	Lys	Val	Leu 495	Gln
Thr	Met	Leu	Asp 500	Thr	Lys	Ala	Thr	Asn 505	Trp	Lys	Glu	Phe	Leu 510	Lys	Glu
Ile	Asp	Glu 515	Lys	Gly	Leu	Asp	Asp 520	Asp	Asp	Leu	Ile	Ile 525	Gly	Leu	Lys
Gly	Lys 530	Glu	Arg	Glu	Leu	Lys 535	Leu	Ala	Gly	Arg	Phe 540	Phe	Ser	Leu	Met
Ser 545	Trp	Lys	Leu	Arg	Glu 550	Tyr	Phe	Val	Ile	Thr 555	Glu	Tyr	Leu	Ile	Lys 560
Thr	His	Phe	Val	Pro 565	Met	Phe	Lys	Gly	Leu 570	Thr	Met	Ala	Asp	Asp 575	Leu
Thr	Ala	Val	Ile 580	Lys	Lys	Met	Leu	Asp 585	Ser	Ser	Ser	Gly	Gln 590	Gly	Leu

Lys	Ser	Tyr 595	Glu	Ala	Ile	Cys	Ile 600	Ala	Asn	His	Ile	Asp 605	Tyr	Glu	Lys
Trp	Asn 610	Asn	His	Gln	Arg	Lys 615	Leu	Ser	Asn	Gly	Pro 620	Val	Phe	Arg	Val
Met 625	Gly	Gln	Phe	Leu	Gly 630	Tyr	Pro	Ser	Leu	Ile 635	Glu	Arg	Thr	His	Glu 640
Phe	Phe	Glu	Lys	Ser 645	Leu	Ile	Tyr	Tyr	Asn 650	Gly	Arg	Pro	Asp	Leu 655	Met
Arg	Val	His	Asn 660	Asn	Thr	Leu	Ile	Asn 665	Ser	Thr	Ser	Gln	Arg 670	Val	Cys
Trp	Gln	Gly 675	Gln	Glu	Gly	Gly	Leu 680	Glu	Gly	Leu	Arg	Gln 685	Lys	Gly	Trp
Thr	Ile 690	Leu	Asn	Leu	Leu	Val 695	Ile	Gln	Arg	Glu	Ala 700	Lys	Ile	Arg	Asn
Thr 705	Ala	Val	Lys	Val	Leu 710	Ala	Gln	Gly	Asp	Asn 715	Gln	Val	Ile	Cys	Thr 720
Gln	Tyr	Lys	Thr	Lys 725	Lys	Ser	Arg	Asn	Val 730	Val	Glu	Leu	Gln	Gly 735	Ala
Leu	Asn	Gln	Met 740	Val	Ser	Asn	Asn	Glu 745	Lys	Ile	Met	Thr	Ala 750	Ile	Lys
Ile	Gly	Thr 755	Gly	Lys	Leu	Gly	Leu 760	Leu	Ile	Asn	Asp	Asp 765	Glu	Thr	Met
Gln	Ser 770	Ala	Asp	Tyr	Leu	Asn 775	Tyr	Gly	Lys	Ile	Pro 780	Ile	Phe	Arg	Gly
Val 785	Ile	Arg	Gly	Leu	Glu 790	Thr	Lys	Arg	Trp	Ser 795	Arg	Val	Thr	Cys	Val 800
Thr	Asn	Asp	Gln	Ile 805	Pro	Thr	Cys	Ala	Asn 810	Ile	Met	Ser	Ser	Val 815	Ser
Thr	Asn	Ala	Leu 820	Thr	Val	Ala	His	Phe 825	Ala	Glu	Asn	Pro	Ile 830	Asn	Ala
Met	Ile	Gln 835	Tyr	Asn	Tyr	Phe	Gly 840	Thr	Phe	Ala	Arg	Leu 845	Leu	Leu	Met
Met	His 850	Asp	Pro	Ala	Leu	Arg 855	Gln	Ser	Leu	Tyr	Glu 860	Val	Gln	Asp	Lys
Ile 865	Pro	Gly	Leu	His	Ser 870	Ser	Thr	Phe	Lys	Tyr 875	Ala	Met	Leu	Tyr	Leu 880
Asp	Pro	Ser	Ile	Gly 885	Gly	Val	Ser	Gly	Met 890	Ser	Leu	Ser	Arg	Phe 895	Leu

.

- Ile Arg Ala Phe Pro Asp Pro Val Thr Glu Ser Leu Ser Phe Trp Arg 900 905 910
- Phe Ile His Val His Ala Arg Ser Glu His Leu Lys Glu Met Ser Ala 915 920 925
- Val Phe Gly Asn Pro Glu Ile Ala Lys Phe Arg Ile Thr His Ile Asp 930 935 940
- Lys Leu Val Glu Asp Pro Thr Ser Leu Asn Ile Ala Met Gly Met Ser 945 950 955 960
- Pro Ala Asn Leu Leu Lys Thr Glu Val Lys Lys Cys Leu Ile Glu Ser 965 970 975
- Arg Gln Thr Ile Arg Asn Gln Val Ile Lys Asp Ala Thr Ile Tyr Leu 980 985 990
- Tyr His Glu Glu Asp Arg Leu Arg Ser Phe Leu Trp Ser Ile Asn Pro 995 1000 1005
- Leu Phe Pro Arg Phe Leu Ser Glu Phe Lys Ser Gly Thr Phe Leu Gly 1010 1015 1020
- Val Ala Asp Gly Leu Ile Ser Leu Phe Gln Asn Ser Arg Thr Ile Arg 1025 1030 1035 1040
- Asn Ser Phe Lys Lys Lys Tyr His Arg Glu Leu Asp Asp Leu Ile Val 1045 1050 1055
- Arg Ser Glu Val Ser Ser Leu Thr His Leu Gly Lys Leu His Leu Arg 1060 1065 1070
- Arg Gly Ser Cys Lys Met Trp Thr Cys Ser Ala Thr His Ala Asp Thr 1075 1080 1085
- Leu Arg Tyr Lys Ser Trp Gly Arg Thr Val Ile Gly Thr Thr Val Pro 1090 1095 1100
- His Pro Leu Glu Met Leu Gly Pro Gln His Arg Lys Glu Thr Pro Cys 1105 1110 1115 1120
- Ala Pro Cys Asn Thr Ser Gly Phe Asn Tyr Val Ser Val His Cys Pro 1125 1130 1135
- Asp Gly Ile His Asp Val Phe Ser Ser Arg Gly Pro Leu Pro Ala Tyr 1140 1145 1150
- Leu Gly Ser Lys Thr Ser Glu Ser Thr Ser Ile Leu Gln Pro Trp Glu 1155 1160 1165
- Arg Glu Ser Lys Val Pro Leu Ile Lys Arg Ala Thr Arg Leu Arg Asp 1170 1175 1180
- Ala Ile Ser Trp Phe Val Glu Pro Asp Ser Lys Leu Ala Met Thr Ile 1185 1190 1195 1200

- Leu Ser Asn Ile His Ser Leu Thr Gly Glu Glu Trp Thr Lys Arg Gln 1205 1210 1215
- His Gly Phe Lys Arg Thr Gly Ser Ala Leu His Arg Phe Ser Thr Ser 1220 1225 1230
- Arg Met Ser His Gly Gly Phe Ala Ser Gln Ser Thr Ala Ala Leu Thr 1235 1240 1245
- Arg Leu Met Ala Thr Thr Asp Thr Met Arg Asp Leu Gly Asp Gln Asn 1250 1255 1260
- Phe Asp Phe Leu Phe Gln Ala Thr Leu Leu Tyr Ala Gln Ile Thr Thr 1265 1270 1275 1280
- Thr Val Ala Arg Asp Gly Trp Ile Thr Ser Cys Thr Asp His Tyr His 1285 1290 1295
- Ile Ala Cys Lys Ser Cys Leu Arg Pro Ile Glu Glu Ile Thr Leu Asp 1300 1305 1310
- Ser Ser Met Asp Tyr Thr Pro Pro Asp Val Ser His Val Leu Lys Thr 1315 1320 1325
- Trp Arg Asn Gly Glu Gly Ser Trp Gly Gln Glu Ile Lys Gln Ile Tyr 1330 1335 1340
- Pro Leu Glu Gly Asn Trp Lys Asn Leu Ala Pro Ala Glu Gln Ser Tyr 1345 1350 1355 1360
- Gln Val Gly Arg Cys Ile Gly Phe Leu Tyr Gly Asp Leu Ala Tyr Arg 1365 1370 1375
- Lys Ser Thr His Ala Glu Asp Ser Ser Leu Phe Pro Leu Ser Ile Gln 1380 1385 1390
- Gly Arg Ile Arg Gly Arg Gly Phe Leu Lys Gly Leu Leu Asp Gly Leu 1395 1400 1405
- Met Arg Ala Ser Cys Cys Gln Val Ile His Arg Arg Ser Leu Ala His 1410 1415 1420
- Leu Lys Arg Pro Ala Asn Ala Val Tyr Gly Gly Leu Ile Tyr Leu Ile 1425 1430 1435 1440
- Asp Lys Leu Ser Val Ser Pro Pro Phe Leu Ser Leu Thr Arg Ser Gly
 1445 1450 1455
- Pro Ile Arg Asp Glu Leu Glu Thr Ile Pro His Lys Ile Pro Thr Ser 1460 1465 1470
- Tyr Pro Thr Ser Asn Arg Asp Met Gly Val Ile Val Arg Asn Tyr Phe 1475 1480 1485
- Lys Tyr Gln Cys Arg Leu Ile Glu Lys Gly Lys Tyr Arg Ser His Tyr 1490 1495 1500

- Ser Gln Leu Trp Leu Phe Ser Asp Val Leu Ser Ile Asp Phe Ile Gly 1505 1510 1515 1520
- Pro Phe Ser Ile Ser Thr Thr Leu Leu Gln Ile Leu Tyr Lys Pro Phe 1525 1530 1535
- Leu Ser Gly Lys Asp Lys Asn Glu Leu Arg Glu Leu Ala Asn Leu Ser 1540 1545 1550
- Ser Leu Leu Arg Ser Gly Glu Gly Trp Glu Asp Ile His Val Lys Phe 1555 1560 1565
- Phe Thr Lys Asp Ile Leu Leu Cys Pro Glu Glu Ile Arg His Ala Cys 1570 1575 1580
- Lys Phe Gly Ile Ala Lys Asp Asn Asn Lys Asp Met Ser Tyr Pro Pro 1585 1590 1595
- Trp Gly Arg Glu Ser Arg Gly Thr Ile Thr Thr Ile Pro Val Tyr Tyr
 1605 1610 1615
- Thr Thr Thr Pro Tyr Pro Lys Met Leu Glu Met Pro Pro Arg Ile Gln 1620 1625 1630
- Asn Pro Leu Ser Gly Ile Arg Leu Gly Gln Leu Pro Thr Gly Ala 1635 1640 1645
- His Tyr Lys Ile Arg Ser Ile Leu His Gly Met Gly Ile His Tyr Arg 1650 1655 1660
- Asp Phe Leu Ser Cys Gly Asp Gly Ser Gly Gly Met Thr Ala Ala Leu 1665 1670 1680
- Leu Arg Glu Asn Val His Ser Arg Gly Ile Phe Asn Ser Leu Leu Glu 1685 1690 1695
- Leu Ser Gly Ser Val Met Arg Gly Ala Ser Pro Glu Pro Pro Ser Ala 1700 1705 1710
- Leu Glu Thr Leu Gly Gly Asp Lys Ser Arg Cys Val Asn Gly Glu Thr 1715 1720 1725
- Cys Trp Glu Tyr Pro Ser Asp Leu Cys Asp Pro Arg Thr Trp Asp Tyr 1730 1735 1740
- Phe Leu Arg Leu Lys Ala Gly Leu Gly Leu Gln Ile Asp Leu Ile Val 1745 1750 1755 1760
- Met Asp Met Glu Val Arg Asp Ser Ser Thr Ser Leu Lys Ile Glu Thr 1765 1770 1775
- Asn Val Arg Asn Tyr Val His Arg Ile Leu Asp Glu Gln Gly Val Leu 1780 1785 1790
- Ile Tyr Lys Thr Tyr Gly Thr Tyr Ile Cys Glu Ser Glu Lys Asn Ala 1795 1800 1805

- Val Thr Ile Leu Gly Pro Met Phe Lys Thr Val Asp Leu Val Gln Thr 1810 1815 1820
- Glu Phe Ser Ser Gln Thr Ser Glu Val Tyr Met Val Cys Lys Gly 1825 1830 1835 1840
- Leu Lys Lys Leu Ile Asp Glu Pro Asn Pro Asp Trp Ser Ser Ile Asn 1845 1850 1855
- Glu Ser Trp Lys Asn Leu Tyr Ala Phe Gln Ser Ser Glu Gln Glu Phe 1860 1865 1870
- Ala Arg Ala Lys Lys Val Ser Thr Tyr Phe Thr Leu Thr Gly Ile Pro 1875 1880 1885
- Ser Gln Phe Ile Pro Asp Pro Phe Val Asn Ile Glu Thr Met Leu Gln 1890 1895 1900
- Ile Phe Gly Val Pro Thr Gly Val Ser His Ala Ala Ala Leu Lys Ser 1905 1910 1915 1920
- Ser Asp Arg Pro Ala Asp Leu Leu Thr Ile Ser Leu Phe Tyr Met Ala 1925 1930 1935
- Ile Ile Ser Tyr Tyr Asn Ile Asn His Ile Arg Val Gly Pro Ile Pro
 1940 1945 1950
- Pro Asn Pro Pro Ser Asp Gly Ile Ala Gln Asn Val Gly Ile Ala Ile 1955 1960 1965
- Thr Gly Ile Ser Phe Trp Leu Ser Leu Met Glu Lys Asp Ile Pro Leu 1970 1975 1980
- Tyr Gln Gln Cys Leu Ala Val Ile Gln Gln Ser Phe Pro Ile Arg Trp 1985 1990 1995 2000
- Glu Ala Val Ser Val Lys Gly Gly Tyr Lys Gln Lys Trp Ser Thr Arg 2005 2010 2015
- Gly Asp Gly Leu Pro Lys Asp Thr Arg Thr Ser Asp Ser Leu Ala Pro 2020 2025 2030
- Ile Gly Asn Trp Ile Arg Ser Leu Glu Leu Val Arg Asn Gln Val Arg 2035 2040 2045
- Leu Asn Pro Phe Asn Glu Ile Leu Phe Asn Gln Leu Cys Arg Thr Val 2050 2055 2060
- Asp Asn His Leu Lys Trp Ser Asn Leu Arg Arg Asn Thr Gly Met Ile 2065 2070 2075 2080
- Glu Trp Ile Asn Arg Arg Ile Ser Lys Glu Asp Arg Ser Ile Leu Met 2085 2090 2095
- Leu Lys Ser Asp Leu His Glu Glu Asn Ser Trp Arg Asp 2100 2105

```
<210> 51
```

<211> 1753

<212> PRT

<213> Vesicular stomatitis virus

<220>

<221> MOD_RES

<222> (1)..(1753)

<223> "Xaa" represents any, other or unknown amino acid

<400> 51

His Trp Gly His Pro Phe Ile Asp Tyr Tyr Ala Gly Leu Glu Lys Leu 1 5 10 15

His Ser Gln Val Thr Xaa Lys Lys Asp Ile Asp Val Ser Tyr Ala Lys 20 25 30

Ala Leu Ala Ser Asp Leu Ala Arg Ile Val Leu Phe Gln Gln Phe Asn 35 40 45

Asp His Xaa Lys Trp Phe Val Asn Gly Asp Leu Leu Pro His Asp His 50 55 60

Pro Phe Lys Ser His Val Lys Glu Asn Thr Trp Pro Thr Ala Ala Gln 65 70 75 80

Val Gln Asp Phe Gly Asp Lys Trp His Glu Leu Pro Leu Ile Lys Cys 85 90 95

Phe Glu Ile Pro Asp Leu Leu Asp Pro Ser Ile Ile Tyr Ser Asp Lys 100 105 110

Ser His Ser Met Asn Arg Ser Glu Val Leu Lys His Val Arg Met Asn 115 120 125

Pro Asn Thr Pro Ile Pro Ser Lys Lys Val Leu Gln Thr Met Leu Asp 130 135 140

Thr Lys Ala Thr Asn Trp Lys Glu Phe Leu Lys Glu Ile Asp Glu Lys 145 150 155 160

Gly Leu Asp Asp Asp Leu Ile Ile Gly Leu Lys Gly Lys Glu Arg 165 170 175

Glu Leu Lys Leu Ala Gly Arg Phe Phe Ser Leu Met Ser Trp Lys Leu 180 185 190

Arg Glu Tyr Phe Val Ile Thr Glu Tyr Leu Ile Lys Thr His Phe Val 195 200 205

Pro Met Phe Lys Gly Leu Thr Met Ala Asp Asp Leu Thr Ala Val Ile 210 215 220

Lys Lys Met Leu Asp Ser Ser Ser Gly Gln Gly Leu Lys Ser Tyr Glu 225 230 235 240

Ala	IIe	Cys	iie	A1a 245	Asn	HIS	iie	Asp	250	GIU	ьуs	Trp	ASI	255	HIS
Gln	Arg	Lys	Leu 260	Ser	Asn	Gly	Pro	Val 265	Phe	Arg	Val	Met	Gly 270	Gln	Phe
Leu	Gly	Tyr 275	Pro	Ser	Leu	Ile	Glu 280	Arg	Thr	His	Glu	Phe 285	Phe	Glu	Lys
Ser	Leu 290		Tyr	Tyr	Asn	Gly 295	Arg	Pro	Asp	Leu	Met 300	Arg	Val	His	Asn
Asn 305	Thr	Leu	Ile	Asn	Ser 310	Thr	Ser	Gln	Arg	Val 315	Cys	Trp	Gln	Gly	Gln 320
Glu	Gly	Gly	Leu	Glu 325	Gly	Leu	Arg	Gln	Lys 330	Gly	Trp	Ser	Ile	Leu 335	Asn
Leu	Leu	Val	Ile 340	Gln	Arg	Glu	Ala	Lys 345	Ile	Arg	Asn	Thr	Ala 350	Val	Lys
Val	Leu	Ala 355	Gln	Gly	Asp	Asn	Gln 360	Val	Ile	Cys	Thr	Gln 365	Tyr	Lys	Thr
Lys	Lys 370	Ser	Arg	Asn	Val	Val 375	Glu	Leu	Gln	Gly	Ala 380	Leu	Asn	Gln	Met
Val 385	Ser	Asn	Asn	Glu	Lys 390	Ile	Met	Thr	Ala	Ile 395	Lys	Ile	Gly	Thr	Gly 400
Lys	Leu	Gly	Leu	Leu 405	Ile	Asn	Asp	Asp	Glu 410	Thr	Met	Gln	Ser	Ala 415	Asp
Tyr	Leu	Asn	Tyr 420	Gly	Lys	Ile	Pro	Ile 425	Phe	Arg	Gly	Val	Ile 430	Arg	Gly
Leu	Glu	Thr 435	Lys	Arg	Trp	Ser	Arg 440	Val	Thr	Cys	Val	Thr 445	Asn	Asp	Gln
Ile	Pro 450	Thr	Cys	Ala	Asn	Ile 455	Met	Ser	Ser	Val	Ser 460	Thr	Asn	Ala	Leu
Thr 465	Val	Ala	His	Phe	Ala 470	Glu	Asn	Pro	Ile	Asn 475	Ala	Met	Ile	Gln	Tyr 480
Asn	Tyr	Phe	Gly	Thr 485	Phe	Ala	Arg	Leu	Leu 490	Leu	Met	Met	His	Asp 495	Pro
Ala	Leu	Arg	Gln 500	Ser	Leu	Tyr	Glu	Val 505	Gln	Asp	Lys	Ile	Pro 510	Gly	Leu
His	Ser	Ser 515	Thr	Phe	Lys	Tyr	Ala 520	Met	Leu	Tyr	Leu	Asp 525	Pro	Ser	Ile
Gly	Gly 530	Val	Ser	Gly	Met	Ser 535	Leu	Ser	Arg	Phe	Leu 540	Ile	Arg	Ala	Phe

Pro 545	Asp	Pro	Val	Thr	Glu 550	Ser	Leu	Ser	Phe	Trp 555	Arg	Phe	Ile	His	Val 560
His	Ala	Arg	Ser	Glu 565	His	Leu	Lys	Glu	Met 570	Ser	Ala	Val	Phe	Gly 575	Asn
Pro	Glu	Ile	Ala 580	Lys	Phe	Arg	Ile	Thr 585	His	Ile	Asp	Lys	Leu 590	Val	Glu
Asp	Pro	Thr 595	Ser	Leu	Asn	Ile	Ala 600	Met	Gly	Met	Ser	Pro 605	Ala	Asn	Leu
Leu	Lys 610	Thr	Glu	Val	Lys	Lys 615	Cys	Leu	Ile	Glu	Ser 620	Arg	Gln	Thr	Ile
Arg 625	Asn	Gln	Val	Ile	Lys 630	Asp	Ala	Thr	Ile	Tyr 635	Leu	Tyr	His	Glu	Glu 640
Asp	Arg	Leu	Arg	Ser 645	Phe	Leu	Trp	Ser	Ile 650	Asn	Pro	Leu	Phe	Pro 655	Arg
Phe	Leu	Ser	Glu 660	Phe	Lys	Ser	Gly	Thr 665	Phe	Leu	Gly	Val	Ala 670	Asp	Gly
Leu	Ile	Ser 675	Leu	Phe	Gln	Asn	Ser 680	Arg	Thr	Ile	Arg	Asn 685	Ser	Phe	Lys
Lys	Lys 690	Tyr	His	Arg	Glu	Leu 695	Asp	Asp	Leu	Ile	Val 700	Arg	Ser	Glu	Val
Ser 705	Ser	Leu	Thr	His	Leu 710	Gly	Lys	Leu	His	Leu 715	Arg	Arg	Gly	Ser	Cys 720
Lys	Met	Trp	Thr	Cys 725	Ser	Ala	Thr	His	Ala 730	Asp	Thr	Leu	Arg	Tyr 735	Lys
Ser	Trp	Gly	Arg 740	Thr	Val	Ile	Gly	Thr 745	Thr	Val	Pro	His	Pro 750	Leu	Glu
Met	Leu	Gly 755	Pro	Gln	His	Arg	Lys 760	Glu	Thr	Pro	Суѕ	Ala 765	Pro	Cys	Asn
Thr	Ser 770	Gly	Phe	Asn	Tyr	Val 775	Ser	Val	His	Cys	Pro 780	Asp	Gly	Ile	His
Asp 785	Val	Phe	Ser	Ser	Arg 790	Gly	Pro	Leu	Pro	Ala 795	Tyr	Leu	Gly	Ser	Lys 800
Thr	Ser	Glu	Ser	Thr 805	Ser	Ile	Leu	Gln	Pro 810	Trp	Glu	Arg	Glu	Ser 815	Lys
Val	Pro	Leu	Ile 820	Lys	Arg	Ala	Thr	Arg 825	Leu	Arg	Asp	Ala	Ile 830	Ser	Trp
Phe	Val	Glu 835	Pro	Asp	Ser	Lys	Leu 840	Ala	Met	Thr	Ile	Leu 845	Ser	Asn	Ile

- His Ser Leu Thr Gly Glu Glu Trp Thr Lys Arg Gln His Gly Phe Lys 850 855 860
- Arg Thr Gly Ser Ala Leu His Arg Phe Ser Thr Ser Arg Met Ser His 865 870 875 880
- Gly Gly Phe Ala Ser Gln Ser Thr Ala Ala Leu Thr Arg Leu Met Ala 885 890 895
- Thr Thr Asp Thr Met Arg Asp Leu Gly Asp Gln Asn Phe Asp Phe Leu 900 905 910
- Phe Gln Ala Thr Leu Leu Tyr Ala Gln Ile Thr Thr Val Ala Arg 915 920 925
- Asp Gly Trp Ile Thr Ser Cys Thr Asp His Tyr His Ile Ala Cys Lys 930 935 940
- Ser Cys Leu Arg Pro Ile Glu Glu Ile Thr Leu Asp Ser Ser Met Asp 945 950 955 960
- Tyr Thr Pro Pro Asp Val Ser His Val Leu Lys Thr Trp Arg Asn Gly 965 970 975
- Glu Gly Ser Trp Gly Gln Glu Ile Lys Gln Ile Tyr Pro Leu Glu Gly 980 985 990
- Asn Trp Lys Asn Leu Ala Pro Ala Glu Gln Ser Tyr Gln Val Gly Arg 995 1000 1005
- Cys Ile Gly Phe Leu Tyr Gly Asp Leu Ala Tyr Arg Lys Ser Thr His 1010 1015 1020
- Ala Glu Asp Ser Ser Leu Phe Pro Leu Ser Ile Gln Gly Arg Ile Arg 1025 1030 1035 1040
- Gly Arg Gly Phe Leu Lys Gly Leu Leu Asp Gly Leu Met Arg Ala Ser 1045 1050 1055
- Cys Cys Gln Val Ile His Arg Arg Ser Leu Ala His Leu Lys Arg Pro 1060 1065 1070
- Ala Asn Ala Val Tyr Gly Gly Leu Ile Tyr Leu Ile Asp Lys Leu Ser 1075 1080 1085
- Val Ser Pro Pro Phe Leu Ser Leu Thr Arg Ser Gly Pro Ile Arg Asp 1090 1095 1100
- Glu Leu Glu Thr Ile Pro His Lys Ile Pro Thr Ser Tyr Pro Thr Ser 1105 1110 1115 1120
- Asn Arg Asp Met Gly Val Ile Val Arg Asn Tyr Phe Lys Tyr Gln Cys 1125 1130 1135
- Arg Leu Ile Glu Lys Gly Lys Tyr Arg Ser His Tyr Ser Gln Leu Trp 1140 1145 1150

- Leu Phe Ser Asp Val Leu Ser Ile Asp Phe Ile Gly Pro Phe Ser Ile 1155 1160 1165
- Ser Thr Thr Leu Leu Gln Ile Leu Tyr Lys Pro Phe Leu Ser Gly Lys 1170 1175 1180
- Asp Lys Asn Glu Leu Arg Glu Leu Ala Asn Leu Ser Ser Leu Leu Arg 1185 1190 1195 1200
- Ser Gly Glu Gly Trp Glu Asp Ile His Val Lys Phe Phe Thr Lys Asp 1205 1210 1215
- Ile Leu Cys Pro Glu Glu Ile Arg His Ala Cys Lys Phe Gly Ile 1220 1225 1230
- Ala Lys Asp Asn Asn Lys Asp Met Ser Tyr Pro Pro Trp Gly Arg Glu 1235 1240 1245
- Ser Arg Gly Thr Ile Thr Thr Ile Pro Val Tyr Tyr Thr Thr Thr Pro 1250 1255 1260
- Tyr Pro Lys Met Leu Glu Met Pro Pro Arg Ile Gln Asn Pro Leu Leu 1265 1270 1275 1280
- Ser Gly Ile Arg Leu Gly Gln Leu Pro Thr Gly Ala His Tyr Lys Ile 1285 1290 1295
- Arg Ser Ile Leu His Gly Met Gly Ile His Tyr Arg Asp Phe Leu Ser 1300 1305 1310
- Cys Gly Asp Gly Ser Gly Gly Met Thr Ala Ala Leu Leu Arg Glu Asn 1315 1320 1325
- Val His Ser Arg Gly Ile Phe Asn Ser Leu Leu Glu Leu Ser Gly Ser 1330 1335 1340
- Val Met Arg Gly Ala Ser Pro Glu Pro Pro Ser Ala Leu Glu Thr Leu 1345 1350 1355 1360
- Gly Gly Asp Lys Ser Arg Cys Val Asn Gly Glu Thr Cys Trp Glu Tyr 1365 1370 1375
- Pro Ser Asp Leu Cys Asp Pro Arg Thr Trp Asp Tyr Phe Leu Arg Leu 1380 1385 1390
- Lys Ala Gly Leu Gly Leu Gln Ile Asp Leu Ile Val Met Asp Met Glu 1395 1400 1405
- Val Arg Asp Ser Ser Thr Ser Leu Lys Ile Glu Thr Asn Val Arg Asn 1410 1415 1420
- Tyr Val His Arg Ile Leu Asp Glu Gln Gly Val Leu Ile Tyr Lys Thr 1425 1430 1435 1440
- Tyr Gly Thr Tyr Ile Cys Glu Ser Glu Lys Asn Ala Val Thr Ile Leu 1445 1450 1455

- Gly Pro Met Phe Lys Thr Val Asp Leu Val Gln Thr Glu Phe Ser Ser 1460 1465 1470
- Ser Gln Thr Ser Glu Val Tyr Met Val Cys Lys Gly Leu Lys Lys Leu 1475 1480 1485
- Ile Asp Glu Pro Asn Pro Asp Trp Ser Ser Ile Asn Glu Ser Trp Lys 1490 1495 1500
- Asn Leu Tyr Ala Phe Gln Ser Ser Glu Gln Glu Phe Ala Arg Ala Lys 1505 1510 1515 1520
- Lys Val Ser Thr Tyr Phe Thr Leu Thr Gly Ile Pro Ser Gln Phe Ile 1525 1530 1535
- Pro Asp Pro Phe Val Asn Ile Glu Thr Met Leu Gln Ile Phe Gly Val 1540 1545 1550
- Pro Thr Gly Val Ser His Ala Ala Ala Leu Lys Ser Ser Asp Arg Pro 1555 1560 1565
- Ala Asp Leu Leu Thr Ile Ser Leu Phe Tyr Met Ala Ile Ile Ser Tyr 1570 1575 1580
- Tyr Asn Ile Asn His Ile Arg Val Gly Pro Ile Pro Pro Asn Pro Pro 1585 1590 1595 1600
- Ser Asp Gly Ile Ala Gln Asn Val Gly Ile Ala Ile Thr Gly Ile Ser 1605 1610 1615
- Phe Trp Leu Ser Leu Met Glu Lys Asp Ile Pro Leu Tyr Gln Gln Cys 1620 1625 1630
- Leu Ala Val Ile Gln Gln Ser Phe Pro Ile Arg Trp Glu Ala Val Ser 1635 1640 1645
- Val Lys Gly Gly Tyr Lys Gln Lys Trp Ser Thr Arg Gly Asp Gly Leu 1650 1655 1660
- Pro Lys Asp Thr Arg Ile Ser Asp Ser Leu Ala Pro Ile Gly Asn Trp 1665 1670 1675 1680
- Ile Arg Ser Leu Glu Leu Val Arg Asn Gln Val Arg Leu Asn Pro Phe 1685 1690 1695
- Asn Glu Ile Leu Phe Asn Gln Leu Cys Arg Thr Val Asp Asn His Leu 1700 1705 1710
- Lys Trp Ser Asn Leu Arg Lys Asn Thr Gly Met Ile Glu Trp Ile Asn 1715 1720 1725
- Arg Arg Ile Ser Lys Glu Asp Arg Ser Ile Leu Met Leu Lys Ser Asp 1730 1735 1740
- Leu His Glu Glu Asn Ser Trp Arg Asp 1745 1750

```
<210> 52
```

<211> 2109

<212> PRT

<213> Vesicular stomatitis virus

<220>

<221> MOD_RES

<222> (1)..(2109)

<223> "Xaa" represents any, other or unknown amino acid

<400> 52

Met Glu Val His Asp Phe Glu Thr Asp Glu Phe Asn Asp Phe Asn Glu

1 1 5 10 15

Asp Asp Tyr Ala Thr Arg Glu Phe Leu Asn Pro Asp Glu Arg Met Thr
20 25 30

Tyr Leu Asn His Ala Asp Tyr Asn Leu Asn Ser Pro Leu Ile Ser Asp 35 40 45

Asp Ile Asp Asn Leu Ile Arg Lys Phe Asn Ser Leu Pro Ile Pro Ser 50 55 60

Met Trp Asp Ser Lys Asn Trp Asp Gly Val Leu Glu Met Leu Thr Ser 65 70 75 80

Cys Gln Ala Asn Pro Ile Ser Thr Ser Gln Met His Lys Trp Met Gly 85 90 95

Ser Trp Leu Met Ser Asp Asn His Asp Ala Ser Gln Gly Tyr Ser Phe
100 105 110

Leu His Glu Val Asp Lys Glu Ala Glu Ile Thr Phe Asp Val Val Glu
115 120 125

Thr Phe Ile Arg Gly Trp Gly Asn Lys Pro Ile Glu Tyr Ile Lys Lys 130 135 140

Glu Arg Trp Thr Asp Ser Phe Lys Ile Leu Ala Tyr Leu Cys Gln Lys 145 150 155 160

Phe Leu Asp Leu His Lys Leu Thr Leu Ile Leu Asn Ala Val Ser Glu 165 170 175

Val Glu Leu Leu Asn Leu Ala Arg Thr Phe Lys Gly Lys Val Arg Arg 180 185 190

Ser Ser His Gly Thr Asn Ile Cys Arg Leu Arg Val Pro Ser Leu Gly 195 200 205

Pro Thr Phe Ile Ser Glu Gly Trp Ala Tyr Phe Lys Lys Leu Asp Ile 210 215 220

Leu Met Asp Arg Asn Phe Leu Leu Met Val Lys Asp Val Ile Ile Gly 225 230 235 240

Arg	Met	GIn	Thr	Val 245	Leu	Ser	Met	Val	Cys 250	Arg	IIe	Asp	Asn	255	Pne
Ser	Glu	Gln	Asp 260	Ile	Phe	Ser	Leu	Leu 265	Asn	Ile	Tyr	Arg	Ile 270	Gly	Asp
Lys	Ile	Val 275	Glu	Arg	Gln	Gly	Asn 280	Phe	Ser	Tyr	Asp	Leu 285	Ile	Lys	Met
Val	Glu 290	Pro	Ile	Суѕ	Asn	Leu 295	Arg	Leu	Met	Lys	Leu 300	Ala	Arg	Glu	Ser
Arg 305	Pro	Leu	Val	Pro	Gln 310	Phe	Pro	His	Phe	Glu 315	Asn	His	Ile	Lys	Thr 320
Ser	Val	Asp	Glu	Gly 325	Ala	Lys	Ile	Asp	Arg 330	Gly	Ile	Arg	Phe	Leu 335	His
Asp	Gln	Ile	Met 340	Ser	Val	Lys	Thr	Val 345	Asp	Leu	Thr	Leu	Val 350	Ile	Tyr
Gly	Ser	Phe 355	Arg	His	Trp	Gly	His 360	Pro	Phe	Ile	Asp	Tyr 365	Tyr	Ala	Gly
Leu	Glu 370	Lys	Leu	His	Ser	Gln 375	Val	Thr	Met	Lys	180 380	Asp	Ile	Asp	Val
Ser 385	Tyr	Ala	Lys	Ala	Leu 390	Ala	Ser	Asp	Leu	Ala 395	Arg	Ile	Val	Leu	Phe 400
				405	-	_	_		410				Asp	415	
Pro	His	Asp	His 420	Pro	Phe	Lys	Ser	His 425	Val	Lys	Glu	Asn	Thr 430	Trp	Pro
		435					440					445	Glu		
	450	_	_			455		_			460		Ser		
Tyr 465	Ser	Asp	Lys	Ser	His 470	Ser	Met	Asn	Arg	Ser 475	Glu	Val	Leu	Lys	His 480
Val	Arg	Met	Asn	Pro 485	Asn	Thr	Pro	Ile	Pro 490	Ser	Lys	Lys	Val	Leu 495	Gln
			500					505					Leu 510		
	_	515	_	_			520		-			525	Gly		_
Gly	Lys 530	Glu	Arg	Glu	Leu	Lys 535	Leu	Ala	Gly	Arg	Phe 540	Phe	Ser	Leu	Met

Ser 545	Trp	Lys	Leu	Arg	Glu 550	Tyr	Phe	Val	Ile	Thr 555	Glu	Tyr	Leu	Ile	Lys 560
Thr	His	Phe	Val	Pro 565	Met	Phe	Lys	Gly	Leu 570	Thr	Met	Ala	Asp	Asp 575	Leu
Thr	Ala	Val	Ile 580	Lys	Lys	Met	Leu	Asp 585	Ser	Ser	Ser	Gly	Gln 590	Gly	Leu
Lys	Ser	Tyr 595	Glu	Ala	Ile	Cys	Ile 600	Ala	Asn	His	Ile	Asp 605	Tyr	Glu	Lys
Trp	Asn 610	Asn	His	Gln	Arg	Lys 615	Leu	Ser	Asn	Gly	Pro 620	Val	Phe	Arg	Val
Met 625	Gly	Gln	Phe	Leu	Gly 630	Tyr	Pro	Ser	Leu	Ile 635	Glu	Arg	Thr	His	Glu 640
Phe	Phe	Glu	Lys	Ser 645	Leu	Ile	Tyr	Tyr	Asn 650	Gly	Arg	Pro	Asp	Leu 655	Met
Arg	Val	His	Asn 660	Asn	Thr	Leu	Ile	Asn 665	Ser	Thr	Ser	Gln	Arg 670	Val	Cys
Trp	Gln	Gly 675	Gln	Glu	Gly	Gly	Leu 680	Glu	Gly	Leu	Arg	Gln 685	Lys	Gly	Trp
Ser	Ile 690	Leu	Asn	Leu	Leu	Val 695	Ile	Gln	Arg	Glu	Ala 700	Lys	Ile	Arg	Asn
Thr 705	Ala	Val	Lys	Val	Leu 710	Ala	Gln	Gly	Asp	Asn 715	Gln	Val	Ile	Cys	Thr 720
Gln	Tyr	Lys	Thr	Lys 725	Lys	Ser	Arg	Asn	Val 730	Val	Glu	Leu	Gln	Gly 735	Ala
Leu	Asn	Gln	Met 740	Val	Ser	Asn	Asn	Glu 745	Lys	Ile	Met	Thr	Ala 750	Ile	Lys
Ile	Gly	Thr 755	Gly	Lys	Leu	Gly	Leu 760	Leu	Ile	Asn	Asp	Asp 765	Glu	Thr	Met
Gln	Ser 770	Ala	Asp	Tyr	Leu	Asn 775	Tyr	Gly	Lys	Ile	Pro 780	Ile	Phe	Arg	Gly
Val 785	Ile	Arg	Gly	Leu	Glu 790	Thr	Lys	Arg	Trp	Ser 795	Arg	Val	Thr	Cys	Val 800
Thr	Asn	Asp	Gln	Ile 805	Pro	Thr	Cys	Ala	Asn 810	Ile	Met	Ser	Ser	Val 815	Ser
Thr	Asn	Ala	Leu 820	Thr	Val	Ala	His	Phe 825	Ala	Glu	Asn	Pro	Ile 830	Asn	Ala
Met	Ile	Gln 835	Tyr	Asn	Tyr	Phe	Gly 840	Thr	Phe	Ala	Arg	Leu 845	Leu	Leu	Met

- Met His Asp Pro Ala Leu Arg Gln Ser Leu Tyr Glu Val Gln Asp Lys 850 855 860
- Ile Pro Gly Leu His Ser Ser Thr Phe Lys Tyr Ala Met Leu Tyr Leu 865 870 875 880
- Asp Pro Ser Ile Gly Gly Val Ser Gly Met Ser Leu Ser Arg Phe Leu 885 890 895
- Ile Arg Ala Phe Pro Asp Pro Val Thr Glu Ser Leu Ser Phe Trp Arg
 900 905 910
- Phe Ile His Val His Ala Arg Ser Glu His Leu Lys Glu Met Ser Ala 915 920 925
- Val Phe Gly Asn Pro Glu Ile Ala Lys Phe Arg Ile Thr His Ile Asp 930 935 940
- Lys Leu Val Glu Asp Pro Thr Ser Leu Asn Ile Ala Met Gly Met Ser 945 950 955 960
- Pro Ala Asn Leu Leu Lys Thr Glu Val Lys Lys Cys Leu Ile Glu Ser 965 970 975
- Arg Gln Thr Ile Arg Asn Gln Val Ile Lys Asp Ala Thr Ile Tyr Leu 980 985 990
- Tyr His Glu Glu Asp Arg Leu Arg Ser Phe Leu Trp Ser Ile Asn Pro 995 1000 1005
- Leu Phe Pro Arg Phe Leu Ser Glu Phe Lys Ser Gly Thr Phe Leu Gly 1010 1015 1020
- Val Ala Asp Gly Leu Ile Ser Leu Phe Gln Asn Ser Arg Thr Ile Arg 1025 1030 1035 1040
- Asn Ser Phe Lys Lys Tyr His Arg Glu Leu Asp Asp Leu Ile Val 1045 1050 1055
- Arg Ser Glu Val Ser Ser Leu Thr His Leu Gly Lys Leu His Leu Arg 1060 1065 1070
- Arg Gly Ser Cys Lys Met Trp Thr Cys Ser Ala Thr His Ala Asp Thr 1075 1080 1085
- Leu Arg Tyr Lys Ser Trp Gly Arg Thr Val Ile Gly Thr Thr Val Pro 1090 1095 1100
- His Pro Leu Glu Met Leu Gly Pro Gln His Arg Lys Glu Thr Pro Cys 1105 1110 1115 1120
- Ala Pro Cys Asn Thr Ser Gly Phe Asn Tyr Val Ser Val His Cys Pro 1125 1130 1135
- Asp Gly Ile His Asp Val Phe Ser Ser Arg Gly Pro Leu Pro Ala Tyr 1140 1145 1150

- Leu Gly Ser Lys Thr Ser Glu Ser Thr Ser Ile Leu Gln Pro Trp Glu 1155 1160 1165
- Arg Glu Ser Lys Val Pro Leu Ile Lys Arg Ala Thr Arg Leu Arg Asp 1170 1175 1180
- Ala Ile Ser Trp Phe Val Glu Pro Asp Ser Lys Leu Ala Met Thr Ile 1185 1190 1195 1200
- Leu Ser Asn Ile His Ser Leu Thr Gly Glu Glu Trp Thr Lys Arg Gln
 1205 1210 1215
- His Gly Phe Lys Arg Thr Gly Ser Ala Leu His Arg Phe Ser Thr Ser 1220 1225 1230
- Arg Met Ser His Gly Gly Phe Ala Ser Gln Ser Thr Ala Ala Leu Thr 1235 1240 1245
- Arg Leu Met Ala Thr Thr Asp Thr Met Arg Asp Leu Gly Asp Gln Asn 1250 1255 1260
- Phe Asp Phe Leu Phe Xaa Ala Thr Leu Leu Tyr Ala Xaa Ile Thr Thr 1265 1270 1275 1280
- Thr Val Ala Arg Asp Gly Trp Ile Thr Ser Cys Thr Asp His Tyr His
 1285 1290 1295
- Ile Ala Cys Lys Ser Cys Leu Arg Pro Ile Glu Glu Ile Thr Leu Asp 1300 1305 1310
- Ser Ser Met Asp Tyr Thr Pro Pro Asp Val Ser His Val Leu Lys Thr 1315 1320 1325
- Trp Arg Asn Gly Glu Gly Ser Trp Gly Gln Glu Ile Lys Gln Ile Tyr 1330 1340
- Pro Leu Glu Gly Asn Trp Lys Asn Leu Ala Pro Ala Glu Gln Ser Tyr 1345 1350 1355 1360
- Gln Val Gly Arg Cys Ile Gly Phe Leu Tyr Gly Asp Leu Ala Tyr Arg 1365 1370 1375
- Lys Ser Thr His Ala Glu Asp Ser Ser Leu Phe Pro Leu Ser Ile Gln
 1380 1385 1390
- Gly Arg Ile Arg Gly Arg Gly Phe Leu Lys Gly Leu Leu Asp Gly Leu 1395 1400 1405
- Met Arg Ala Ser Cys Cys Gln Val Ile His Arg Arg Ser Leu Ala His 1410 1415 1420
- Leu Lys Arg Pro Ala Asn Ala Val Tyr Gly Gly Leu Ile Tyr Leu Ile 1425 1430 1435 1440
- Asp Lys Leu Ser Val Ser Pro Pro Phe Leu Ser Leu Thr Arg Ser Gly
 1445 1450 1455

- Pro Ile Arg Asp Glu Leu Glu Thr Ile Pro His Lys Ile Pro Thr Ser 1460 1465 1470
- Tyr Pro Thr Ser Asn Arg Asp Met Gly Val Ile Val Arg Asn Tyr Phe 1475 1480 1485
- Lys Tyr Gln Cys Arg Leu Ile Glu Lys Gly Lys Tyr Arg Ser His Tyr 1490 1495 1500
- Ser Gln Leu Trp Leu Phe Ser Asp Val Leu Ser Ile Asp Phe Ile Gly 1505 1510 1515 1520
- Pro Phe Ser Ile Ser Thr Thr Leu Leu Gln Ile Leu Tyr Lys Pro Phe 1525 1530 1535
- Leu Ser Gly Lys Asp Lys Asn Glu Leu Arg Glu Leu Ala Asn Leu Ser 1540 1545 1550
- Ser Leu Leu Arg Ser Gly Glu Gly Trp Glu Asp Ile His Val Lys Phe 1555 1560 1565
- Phe Thr Lys Asp Ile Leu Cys Pro Glu Glu Ile Arg His Ala Cys 1570 1575 1580
- Lys Phe Gly Ile Ala Lys Asp Asn Asn Lys Asp Met Ser Tyr Pro Pro 1585 1590 1595 1600
- Trp Gly Arg Glu Ser Arg Gly Thr Ile Thr Thr Ile Pro Val Tyr Tyr
 1605 1610 1615
- Thr Thr Pro Tyr Pro Lys Met Leu Glu Met Pro Pro Arg Ile Gln
 1620 1625 1630
- Asn Pro Leu Ser Gly Ile Arg Leu Gly Gln Leu Pro Thr Gly Ala 1635 1640 1645
- His Tyr Lys Ile Arg Ser Ile Leu His Gly Met Gly Ile His Tyr Arg 1650 1655 1660
- Asp Phe Leu Ser Cys Gly Asp Gly Ser Gly Gly Met Thr Ala Ala Leu 1665 1670 1680
- Leu Arg Glu Asn Val His Ser Arg Gly Ile Phe Asn Ser Leu Leu Glu 1685 1690 1695
- Leu Ser Gly Ser Val Met Arg Gly Ala Ser Pro Glu Pro Pro Ser Ala 1700 1705 1710
- Leu Glu Thr Leu Gly Gly Asp Lys Ser Arg Cys Val Asn Gly Glu Thr 1715 1720 1725
- Cys Trp Glu Tyr Pro Ser Asp Leu Cys Asp Pro Arg Thr Trp Asp Tyr 1730 1735 1740
- Phe Leu Arg Leu Lys Ala Gly Leu Gly Leu Gln Ile Asp Leu Ile Val 1745 1750 1755 1760

- Met Asp Met Glu Val Arg Asp Ser Ser Thr Ser Leu Lys Ile Glu Thr 1765 1770 1775
- Asn Val Arg Asn Tyr Val His Arg Ile Leu Asp Glu Gln Gly Val Leu 1780 1785 1790
- Ile Tyr Lys Thr Tyr Gly Thr Tyr Ile Cys Glu Ser Glu Lys Asn Ala 1795 1800 1805
- Val Thr Ile Leu Gly Pro Met Phe Lys Thr Val Asp Leu Val Gln Thr 1810 1815 1820
- Glu Phe Ser Ser Ser Gln Thr Ser Glu Val Tyr Met Val Cys Lys Gly 1825 1830 1835 1840
- Leu Lys Lys Leu Ile Asp Glu Pro Asn Pro Asp Trp Ser Ser Ile Asn 1845 1850 1855
- Glu Ser Trp Lys Asn Leu Tyr Ala Phe Gln Ser Ser Glu Gln Glu Phe 1860 1865 1870
- Ala Arg Ala Lys Lys Val Ser Thr Tyr Phe Thr Leu Thr Gly Ile Pro 1875 1880 1885
- Ser Gln Phe Ile Pro Asp Pro Phe Val Asn Ile Glu Thr Met Leu Gln 1890 1895 1900
- Ile Phe Gly Val Pro Thr Gly Val Ser His Ala Ala Ala Leu Lys Ser 1905 1910 1915 1920
- Ser Asp Arg Pro Ala Asp Leu Leu Thr Ile Ser Leu Phe Tyr Met Ala 1925 1930 1935
- Ile Ile Ser Tyr Tyr Asn Ile Asn His Ile Arg Val Gly Pro Ile Pro
 1940 1945 1950
- Pro Asn Pro Pro Ser Asp Gly Ile Ala Gln Asn Val Gly Ile Ala Ile 1955 1960 1965
- Thr Gly Ile Ser Phe Trp Leu Ser Leu Met Glu Lys Asp Ile Pro Leu 1970 1975 1980
- Tyr Gln Gln Cys Leu Ala Val Ile Gln Gln Ser Phe Pro Ile Arg Trp 1985 1990 1995 2000
- Glu Ala Val Ser Val Lys Gly Gly Tyr Lys Gln Lys Trp Ser Thr Arg 2005 2010 2015
- Gly Asp Gly Leu Pro Lys Asp Thr Arg Ile Ser Asp Ser Leu Ala Pro 2020 2025 2030
- Ile Gly Asn Trp Ile Arg Ser Leu Glu Leu Val Arg Asn Gln Val Arg 2035 2040 2045
- Leu Asn Pro Phe Asn Glu Ile Leu Phe Asn Gln Leu Cys Arg Thr Val 2050 2055 2060

Asp Asn His Leu Lys Trp Ser Asn Leu Arg Lys Asn Thr Gly Met Ile 2065 2070 2075 2080

Glu Trp Ile Asn Arg Arg Ile Ser Lys Glu Asp Arg Ser Ile Leu Met $2085 \hspace{1cm} 2090 \hspace{1cm} 2095$

Leu Lys Ser Asp Leu His Glu Glu Asn Ser Trp Arg Asp 2100 2105